Computer-Aided Diagnosis System for Retinal Diseases in Medical Imaging

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Abstract — Bordering on important domains – engineering, medicine and informatics, involving not only knowledge regarding the biosystems structure and functionality, but also knowledge and skills from technical and IT systems, the paper presents the results of some researches on biological human visual structures concerning the diagnosis of visual diseases, namely the macular ones. This highly important transdisciplinary topic combines aspects from biosystems (human visual system), image acquisition and processing (medical imaging), artificial intelligence techniques (neural networks) and information management (databases). Starting from classical or digital retina images, using neural networks image recognition algorithms, the Computer-Aided Diagnosis (CADx) system identifies macular diseases with high precision. Images are stored in databases together with patient personal details and treatments and diagnosis information. The software includes image processing modules, databases and artificial neural networks that can be trained for recognizing images of new diseases or for improving sensitivity and specificity of the system. The Computer-Aided Diagnosis reduces the doctor's level of incertitude regarding some diseases, improves the initial and evolutional identification precision of disease, allows monitoring the health status of the patient during new treatment methods, stores images in digital format and generates diagnoses databases that can be explored in research, medical practice and specialized teaching. Using such a system increases the quality and accessibility of medical services.

Key-Words: — computer-aided diagnosis, neural networks, image processing, ophthalmology, macula, retina

1 Introduction

During last years, the diagnosis of human visual system diseases had an amazing progress, especially due to technological innovations and exponential development in the IT domain. Taking into account the diversity and complexity of the eye functions, a huge number of equipments, devices, methods and algorithms for diagnosis were developed. Sometimes the doctor can identify certain diseases after a visual analysis of an image representing the affected area. However, in a large number of cases, the diagnosis is not successful, due to different factors as, for example, reduced experience, fatigue, diversity of shapes and texture, similarity, poor image quality and so on. In these situations, a second opinion, coming from another expert or better from a computer-aided diagnosis system, is very important and useful.

From the multitude of the human visual system diseases, those that affect the inside back nervous layer of the eye, namely retina, can be identified by image recognition in a so-called retinal diseases medical imaging process. From the retinal diseases, the macular ones are of the worst.

Macula is a biological structure from the human visual system, responsible for high acuity vision, color vision and central vision. Consisting from a small sensitive area in the central part of retina, macula allows focusing central vision in the eye and controls people ability to read, drive a car, recognize faces or colors and see objects in fine detail [1].

There are different eye conditions in which macula can be damaged, often causing loss of central vision (Fig.1). Macular diseases, especially AMD (Age-Related Macular Degeneration) are an increasing problem worldwide, because they are irreversible, so it is very important that the process to be early identified and stopped. Macular degeneration is caused by the deterioration of the central portion of the retina, the inside back layer of the eye that records the images we see and sends them via the optic nerve from the eye to the brain.



Fig.1. The influence of AMD on human vision [2]: a - normal vision, b – central vision loss due to AMD

Such problems may occur at a higher age, but in the last years, they were observed at the second part of life (40 - 60 years old people).

Current treatment options can delay progression and research continues into ways of reversing retinal damage.

2 Problem Formulation

Macular diseases have a great variety of shapes and textures and sometimes they are hard to be identified and recognized even by experienced doctors. In these situations, a Computer-Aided Diagnosis (CADx) system becomes a very useful tool not only for doctors, but also for researchers. That is why we are trying to develop and optimize such a system based on intelligent image classification/recognition algorithms.

It is very important to make distinction between Computer-Aided Detection (CAD) and Computer-Aided Diagnosis (CADx) [3]. While the first method supposes only to detect and localize the interest areas of a possible disease, the second one obtaining analysis, allows area diagnostic classification or disease recognition with high precision. Regarding performance, a CAD system is characterized by *sensitivity*, representing the percent of images correctly detected as having potential diseases, while a CADx system is characterized by specificity, as the percent of diseases images correctly recognized.

The Computer-Aided Diagnosis (CADx) system for macular diseases is a very useful tool for doctors and researchers due to the following advantages:

- A diagnosis recognized by the computer will reduce the doctor's level of incertitude;
- It offers accuracy, consistence and a high confidence coefficient in results interpreting;

- It allows monitoring the health status of the patient during new treatment methods;
- The diagnosis identification time is significantly reduced;
- The diagnosis can be stored in a digital format (an image file for the retina photo and a medical record in a specific patient database);
- The image can be processed and analyzed on-line or off-line;
- The system generates a diagnoses database useful for researchers, for medical practice and for specialized teaching purposes;
- A database with patient personal and care information can be developed, having local or remote secured access;
- The long treatment costs, the supplementary tests and the number of surgical intervention are significantly reduced;
- The quality and accessibility of medical services are increased.

2.1 The structure of the CADx system

The CADx system has to analyze the image of the patient's macula and recognize it, offering a response that contains four possible diagnoses, ordered by a level of certitude. Initially, the system was trained to recognize the normal macula and a certain set of diagnoses, but it can be further trained, from software, to learn more.

The structure of the CADx process is presented in Fig.2.

The image of the retina is acquired using a retinal investigation camera and transferred to computer as data input for the designed software. An image processing module is used for different image quality improvements or geometrical transformations. A certain vector of image features is then automatically extracted and the obtained values are applied to the inputs of the artificial network (ANN) neural used for image classification/recognition [4]. The ANN generates a response that represents a set of four possible diagnoses, ordered by the level of certitude. In the end, the doctor can establish the final diagnosis after a discussion with the patient, taking into account personal and heredo-collateral antecedents and others aspects considered important. The results are stored in a database, as a patient medical record, together with personal information and recommended treatment.



Fig.2. The structure of the CADx process

2.2 The knowledge stage in the domain

At the international level, the analysis of the present stage showed an extremely important interest, the computer-aided diagnosis based o image recognition being one of the major research topics within the framework of the medical imaging [3], [5]. This interest is motivated by the care shown towards the people's health state, together with the advantages of CADx mentioned above. A minute analysis of the specialty literature stated the following:

- In the field of medical imaging, there have been carried out numberless research studies concerning the computer-aided diagnosis of breast cancer [6], of the pulmonary nodules from breast X-ray pictures [7] and CAT SCANs [8], [9], of the cerebral aneurysms [10], [11], some areas being better covered, other areas less covered, just like the case of retinal imaging;
- In the field of retinal imaging, there is a series of achievements, but because of the great range of diagnosis diversity, the solved issues have in view only a selective treatment of their identification [12]. There are research studies concerning the glaucoma diagnosis in the eye fundus images [13], iris recognition, white spots identification and intensive bleedings at the

retina level, automated detection of diabetic retinopathy damaging [14], but there are no final results having in view a group of diseases;

- The image processing and analysis in this field has generally in view the use of image segmentation operations (of the optical disc, of the macula, vascular arch, macular drusen deposits, microaneurysms and exudates in diabetic retinopathy) [15], contrast enhancement [16], filter application [17];
- The frequently used images classification/recognition methods and techniques use linear or superior order classifiers [18], decision trees, machine learning [19] and most frequently neural networks [20], [21], [22], [23].

3 Problem Solution

In order to solve the computer-aided diagnosis problem for macular diseases and for optimizing the solution, a software package has been designed. Specific image processing module, database module and neural networks module were developed in Matlab®.

A description of the system components is presented forwards, focusing on the most important aspects from the diagnosis process.

3.1 Image acquisition

Retina's image can be acquired directly from a retinal investigation camera that offers a digital image or a classical one on color-reversal film. These last types of pictures have to be processed on a photo processing machine and then digitized using a scanner. The system is ready to work with any image of the retina, in digital format. The file format of the image used for recognition, in this software release, has to be a *jpg* one.

3.2 Image processing

Having the images on the computer screen, the doctor can observe details and process them, in order to emphasize the elements that may identify the diagnosis.

For doing this, a Graphical User Interface (GUI) developed in Matlab by the authors, offers access to an image processing module. It contains options for processing one image at a time (Fig.3) or for

comparing two images, representing for example two stages of a disease of the same patient.



Fig.3. Main window for the image processing module

Comparison can be made from a visual point, as well as from a mathematical point of view – histograms and values of the image features.

Different options are available for user (Fig.3): opening a file, saving it in the same or other format, viewing the original image, the Red (R), Green (G) and Blue (B) components, transforming from RGB to Gray. Histograms for R, G and B components can be computed and plotted for a number of maximum 255 bins. User can zoom, pan and rotate the image. Image details are examined using *Image Tools* facility (Fig.4). Options as *Vertical Zoom, Horizontal Zoom, Unconstrained Zoom, Vertical Pan, Horizontal Pan, Unconstrained Pan* and *Reset* to Original View are available. The image can be horizontal or vertical mirrored.

The contrast can be enhanced using histogram equalization. Doctor can select and crop an interest area of image and save it.



Fig.4. Image Tools option

All of these tools are available for emphasizing some image characteristics so that the diagnostic to be easily recognized.

The possibilities for viewing and editing an image are very useful for user, especially for doctor, but they were very important in the design phase of the Computer-Aided Diagnosis system. They gave the possibility to generate a database with images of diagnoses used for training an artificial neural network in the image recognition process. Another database was used for testing the network.

In the design phase of the system, it was possible, having all these tools, to prepare images so that to see if a rotated image or a mirrored one can be correctly recognized, if it is better to use gray images or RGB images or contrast enhanced images for neural network inputs.

Another important option available in the image processing module is the "*Two Images Comparison*".

Two image files can be opened and analyzed together, visual or using normalized histograms or values that represent features. User can zoom and pan images and plotted histograms.

3.3 Image analysis and features extraction

Macular diseases images are characterized by a large diversity of shapes and structures, so it is more helpful to use for image classification/recognition, texture descriptors instead of shape or boundary descriptors. That is why a set of six values that describe a region by quantifying its texture content were used as image features [24]. The six descriptors are based on statistical properties of the intensity histogram, namely on statistical moments and they are: *mean* – a measure of average intensity, *standard deviation* – a measure of average contrast, *smoothness* of the intensity in that region, *third moment* – a measure of skewness of a histogram, *uniformity* and *entropy*.

The system has to recognize images even if they are rotated, mirrored or scaled. Regarding this aspect, a set of seven 2D moment invariants, insensitive to translation, scale, mirroring and rotation were used.

Special user functions were developed in Matlab for automatically computing the values above [25], [26]. The set of seven 2D moment invariants are computed using the equations below.

For a digital image having a f(x,y) intensity distribution, the two dimensional moment of order p+q is defined as [24]

$$m_{pq} = \sum_{x} \sum_{y} x^{p} y^{q} f(x, y), \qquad (1)$$

for p, q = 0, 1, 2,... and x, y are the values of image coordinates.

These moments in general are not invariant to any distortions and therefore the corresponding central moments are defined as

$$\mu_{pq} = \sum_{x} \sum_{y} \left(x - \overline{x} \right)^p \left(y - \overline{y} \right)^q f(x, y), \tag{2}$$

where we denoted

$$\bar{x} = \frac{m_{10}}{m_{00}}$$
 and $\bar{y} = \frac{m_{01}}{m_{00}}$. (3)

The central moments are known to be invariant under translation. It can be demonstrated that the first four orders of central moments from equation (2) can be expressed in terms of the ordinary moments defined in equation (1).

$$\mu_{00} = m_{00} ,$$

$$\mu_{10} = \mu_{01} = 0 ,$$

$$\mu_{11} = m_{11} - \frac{m_{10}m_{01}}{m_{00}} ,$$

$$\mu_{20} = m_{20} - \frac{m_{10}^2}{m_{00}} ,$$

$$\mu_{02} = m_{02} - \frac{m_{01}^2}{m_{00}} ,$$

$$\mu_{12} = m_{12} - m_{02}\bar{x} - 2m_{11}\bar{y} + 2m_{10}\bar{y}^2 ,$$

$$\mu_{21} = m_{21} - m_{20}\bar{y} - 2m_{11}\bar{x} + 2m_{01}\bar{x}^2 ,$$

$$\mu_{03} = m_{03} - 3m_{02}\bar{y} + 2m_{01}\bar{y}^2 .$$

(4)

Often it is desirable to normalize the moments with respect to size. This may be accomplished by using the area, μ_{00} .

The normalized central moment of order (p + q) is defined as

$$\eta_{pq} = \frac{\mu_{pq}}{\mu_{00}^{\gamma}},$$
(5)
for $p, q = 0, 1, 2, ...$
and
 $\gamma = \frac{p+q}{2} + 1,$ (6)

for p + q = 2, 3, ...

From above equations, a set of seven **2D moment invariants**, insensitive to translation, change, scale, mirroring and rotation can be derived:

$$\begin{split} \phi_{1} &= \eta_{20} + \eta_{02} ,\\ \phi_{2} &= (\eta_{20} - \eta_{02})^{2} + 4\eta_{11}^{2} ,\\ \phi_{3} &= (\eta_{30} - 3\eta_{12})^{2} + (3\eta_{21} - \eta_{03})^{2} ,\\ \phi_{4} &= (\eta_{30} + \eta_{12})^{2} + (\eta_{21} + \eta_{03})^{2} ,\\ \phi_{5} &= (\eta_{30} - 3\eta_{12})(\eta_{30} + \eta_{12}) \cdot \\ \cdot \left[(\eta_{30} + \eta_{12})^{2} - 3(\eta_{21} + \eta_{03})^{2} \right] + \\ &+ (3\eta_{21} - \eta_{03})(\eta_{21} + \eta_{03}) \cdot \\ \cdot \left[3(\eta_{30} + \eta_{12})^{2} - (\eta_{21} + \eta_{03})^{2} \right] + \\ &+ 4\eta_{11}(\eta_{30} + \eta_{12})(\eta_{21} + \eta_{03}) + \\ \phi_{7} &= (3\eta_{21} - \eta_{03})(\eta_{30} + \eta_{12}) \cdot \\ \cdot \left[(\eta_{30} + \eta_{12})^{2} - 3(\eta_{21} + \eta_{03})^{2} \right] + \\ &+ (3\eta_{12} - \eta_{30})(\eta_{21} + \eta_{03}) \cdot \\ \cdot \left[3(\eta_{30} + \eta_{12})^{2} - (\eta_{21} + \eta_{03})^{2} \right] + \\ &+ (3\eta_{12} - \eta_{30})(\eta_{21} + \eta_{03}) \cdot \\ \cdot \left[3(\eta_{30} + \eta_{12})^{2} - (\eta_{21} + \eta_{03})^{2} \right] + \\ \end{split}$$

The other six values representing image's features are also computed with the GUI [26]. An important approach for describing a region is to quantify its texture content. A frequently used method for texture analysis is based on statistical properties of the intensity histogram.

Let us consider z_i a discrete random variable that corresponds to the intensity levels of an image and let us denote $p(z_i)$ the corresponding normalized histogram, with i = 0, 1, 2, ..., L-1 and L – the number of possible intensity values.

One of the principal approaches for describing the shape of a histogram is via its *central moments* (also called *moments about the mean*) [24], which are defined as

$$\mu_n = \sum_{i=0}^{L-1} (z_i - m)^n p(z_i), \qquad (8)$$

where *n* is the moment order and *m* is the *mean*:

$$m = \sum_{i=0}^{L-1} z_i p(z_i).$$
 (9)

The *mean* (*m*), a measure of average intensity, is the **first image feature** in a vector of such values. Because the histogram is assumed to be normalized, the sum of its all components is 1.

The second moment is called the *variance*,

$$\mu_2 = \sum_{i=0}^{L-1} (z_i - m)^2 p(z_i).$$
(10)

Standard deviation (σ), a measure of average contrast, is used also as an image feature,

$$\sigma = \sqrt{\mu_2(z)} = \sqrt{\sigma^2} . \tag{11}$$

The third image feature is the *smoothness* (*R*). It measures the relative smoothness of the intensity in a region. In practice, the variance is normalized to the range [0, 1] by dividing it by $(L-1)^2$,

$$R = 1 - 1/(1 + \sigma^2).$$
(12)

The fourth image feature is the *third order moment* (μ_3). It measures the skewness of a histogram. If the histogram is symmetric than $\mu_3 = 0$, if it is skewed to the right of the mean than $\mu_3 > 0$, and if it is skewed to the left of the mean than $\mu_3 < 0$. Values of this measure are brought into a range of values comparable to the other five measures by dividing μ_3 by $(L-1)^2$,

$$\mu_3 = \sum_{i=0}^{L-1} (z_i - m)^3 p(z_i).$$
(13)

The fifth image feature is *uniformity* (U). It is maximum when all the gray levels are equal (maximally uniform) and decreases from there,

$$U = \sum_{i=0}^{L-1} p^2(z_i).$$
 (14)

The last value in our vector of features is *entropy* (*e*), a measure of randomness,

$$e = \sum_{i=0}^{L-1} p(z_i) \log_2 p(z_i).$$
(15)

All of these values are computed in Matlab® with a user-defined function. In the image processing module of the Graphical User Interface developed for diagnostic recognition, the option *"Two Images Comparison"* allows computing and

comparing them. The computed values are scaled by the vector [1 1 10000 10000 10000 1], so that to work with appropriate ranges.

Using the GUI, it was made a comparison between images using R, G, B components, gray image and the corresponding multidimensional one [26]. It was observed that differences between 2D moment invariants values are not significant, excepting the multidimensional image. It is normal because the multidimensional image is obtained by concatenating the 3 two dimensional arrays (of RGB image) in a 2D array. The rotated multidimensional image is made by 3 two dimensional images rotated before, not by rotating directly the entire multidimensional image. So it is better to use the RGB image or the gray one when extracting features, instead of multidimensional images.

3.4 Artificial neural network for image recognition

An image is made by pixels. Depending on the image format and knowing that each pixel has associated a certain number of values, the total number of values can be a huge one. It is impossible to associate a neuron for each value in the neural network structure. It is almost impossible to find two identical retina images at the pixel level. That is why for the image recognition with artificial neural networks we have to use at inputs a vector of values extracted from image features.

A total connected multilayer perceptron network was chosen for diagnosis recognition, knowing that it offers good results in such type of applications [27]. Steps as configuring, training, testing and optimizing ANN, were followed in the design phase of CADx system (Fig.5), but they are also available in the final version of the software, in a so called administrator mode, for improving the system performance. All these steps are executed using another GUI designed also in Matlab® [28].

For training the ANN, it is necessary to have a training set consisting in different images of macular diseases (including also multiple different images of the same diagnosis). A special database module was developed, useful not only for storing images, but for storing features of the images [29].

The name and path of the image file, the name, number and associated vector for diagnosis, histograms computed for a certain number of bins, features values for the 2D moment invariants and texture descriptors for each of the R, G, B color components or for gray type images, are information stored in the fields of the database (Fig.6).



Fig.5. Image recognition process using ANN

All these values can be computed for each image at a time or automatically for the entire database. New records can be appended and the existing ones can be deleted or modified (Fig.7). Images can be browsed one by one. If an image is an area cropped from another one, it is possible to store in the database information about both of them (very useful option because only macular area is examined by the system, while it is important to see the global image of retina to which the part was associated).

The training set is generated from such a database that contains information about macular diseases images [29].

Field 🔺	Value
Denumire	'Angiopatie hipertensiva'
FisImagine	D:\MACULA TEST\Imagini Complete Color\Angiopatie hipertensiva - part.jpg'
Nr_clase	10
HistoR	[0 0.299 0.5981 0.3738 0.1869 0.299 494.92 445.84 57.491 0]
HistoG	[0 1.1962 0.299 205.41 792.84 0.2617 0 0 0 0]
HistoB	[455.07 182.49 361.69 0.7476 0 0 0 0 0 0]
Obs	
MomentR	[7.0064 17.906 27.536 27.437 55.053 36.413 55.664]
MomentG	[6.4068 16.664 26.778 27.839 55.411 36.178 55.68]
MomentB	[5.5686 13.216 20.163 20.734 42.682 27.865 41.328]
HistoGri	
MomentGri	
DescrR	[187.18 15.594 37.259 -305.19 199.16 5.7933]
DescrG	[102.95 5.939 5.4214 -82.075 611.19 4.381]
DescrB	[26.285 24.413 90.82 366.68 710.04 5.0834]
DescrGri	
Nr_diag	1
Vect_diag	<1x14 double>
FisImagine_ini	D:WACULA TEST\Imagini Complete Color\Angiopatie hipertensiva.jpg'
Descr_scara	[1 1 10000 10000 10000 1]
Diagnostic ×	Diagnostic(1,1) × Diagnostic(1,2) × Diagnostic(1,1).Vect_diag × Diagnostic(1,4) ×



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D:WACU	LA TESTVimagini Complete Color/Drusen macular - part.jpg	Stergere Momente
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D:WACU	LA TESTVimagini Complete Color/Drusen macular.jpg	
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Rosu	[0 0 0 0 0 0 19.9564 560.8623 419.1813]	
	[0 0 0.0964 28.074 348.7843 509.689 112.4308 0.9255 0 0]	
Verde	[994.9289 5.0711 0 0 0 0 0 0 0 0]	
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Fig.7. Main window for the diagnoses database

The number of values representing features will determine the number of inputs in the neural network and it can be selected from the main window of the neural network module (Fig.8). Using RGB images and seven 2D moment invariants plus six texture descriptors, the number of inputs was 39 (13 for each color component).

The number of diagnostics from the database will determine the number of outputs. The structure of the network can be modified by selecting one or two hidden layers and choosing the number of neurons in each layer. The transfer function can be selected from a list that contains *tansig* - hyperbolic tangent sigmoid transfer function, *logsig* - log sigmoid transfer function and *purelin* - linear transfer function.



Fig.8. Main window for ANN configuring, training, testing and optimizing

Because the teacher output vector is composed from 0 and 1 values (1 for marking the correct diagnostic), *logsig* and *tansig* functions will be preferred considering the interval in which they are taking values.

The training parameters and the stopping conditions can be also modified. Learning is a supervised one.

After a large number of successful tests, the optimization stage of the recognition process was started. We have to find out what structure of the neural network, training algorithms and parameters offer the best results.

Optimization criteria take into account three elements: maximizing the number of classified images (first priority – main objective), minimizing the training time (second priority) and minimizing the square mean error (third priority).

Restrictions refer to: the ANN type - total connected feedforward multilayer perceptron network; image database (training set) - 25 RGB images, representing 15 diagnostics; ANN architecture - 39–X–15 with 3 layers of neurons, where X is the variable number of neurons in the hidden layer; training process - 15 cycles with 30.000 epochs each, results being stored at each 1000 epochs.

Three optimization parameters were taken into account: the training function, the transfer function and the ANN architecture, namely the number of neurons in the hidden layer.

Twelve training functions were pretested, six of them being finally used for analysis: *traingd* -Gradient descent backpropagation, *traingdm* -Gradient descent with momentum backpropagation, *traingda* - Gradient descent with adaptive learning rate backpropagation, *traingdx* - Gradient descent with momentum and adaptive learning rate backpropagation, *trainrp* - Resilient backpropagation and *trainscg* - Scaled conjugate gradient backpropagation.

Three transfer function were tested: *tansig* – bipolar sigmoid, *logsig* – unipolar sigmoid, *purelin* – linear function.

Regarding network architecture, three different structures were used, with 10, 39, 78 neurons in the hidden layer.

The analysis can be done manually, gradually, choosing a certain network structure, initializing weights, storing initial weights for using the same values for all the training functions, starting the training process for a certain learning algorithm, storing results during and at the end of the training process, going back to train the network with other function and other structure and so on. This type of analysis is not so easy to be done because of long training time and long processing time of the results. That is why we have developed specific commands for automatically training, storing and processing of the results for each training strategy. Finally, the results are averaged for the number of training cycles.

In Table 1 is presented, for example, the average square mean error for an ANN with a 39-78-15 structure, trained with three different algorithms.

Regarding the training function, best results were obtained for the *trainscg* - scaled conjugate gradient backpropagation algorithm [30]. Taking into account the restrictions mentioned above, after 10.000 training epochs, all 25 images from the database with 15 diagnostics were classified and square mean error is the smallest (Fig.9, Fig.10 - graphs are automatically generated from the software).

Even the training time for this function was higher than those obtained for the other tested functions, there are some methods to eliminate this disadvantage (cumulative training [29]).

All the tests for the training algorithm counted more than 200 training cycles, each of them having 30.000 to 80.000 epochs.

Epochs	Training function				
$[x10^3]$	traingd	traingdm	trainscg		
1	0.091106	0.095071	0.044864		
2	0.075433	0.077289	0.029086		
3	0.065253	0.070717	0.019804		
4	0.057608	0.063157	0.013827		
5	0.052207	0.059571	0.010579		
6	0.048520	0.057414	0.007957		
7	0.046636	0.055822	0.005169		
8	0.044688	0.054163	0.003788		
9	0.043009	0.052996	0.002778		
10	0.041746	0.052050	0.002022		
11	0.040647	0.051241	0.001498		
12	0.039635	0.050446	0.001121		
13	0.038733	0.049565	0.000857		
14	0.037951	0.048906	0.000641		
15	0.037280	0.048252	0.000484		
16	0.036765	0.047672	0.000371		
17	0.036307	0.047196	0.000287		
18	0.036205	0.046791	0.000227		
19	0.035512	0.046402	0.000180		
20	0.034884	0.046031	0.000145		
21	0.035085	0.045602	0.000116		
22	0.034276	0.045203	0.000090		
23	0.033810	0.044834	0.000072		
24	0.033245	0.044472	0.000058		
25	0.032845	0.044129	0.000047		
26	0.032403	0.043726	0.000038		
27	0.031804	0.040679	0.000030		
28	0.031161	0.039840	0.000025		
29	0.030698	0.039499	0.000020		
30	0.030339	0.039221	0.000016		

 Table 1. The average square mean error



Fig.9. Square mean error distribution for 15 training cycles, for 3 training functions

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Regarding the transfer function, best results were obtained for the *tansig* - hyperbolic tangent sigmoid transfer function, even the square mean error is not the smallest. There were no significant differences at the training time for the tested transfer functions, but the mean number of classified images was better for *tansig* and for a bigger number of neurons in the hidden layer (Fig.11).

From the structure of the network point of view, three different configuration were tested, with 10, 39, 78 neurons in the hidden layer. The higher the number of neurons, the higher the time for a training cycle, but the diagnostics should be classified in fewer cycles. Best results for the square mean error and number of classified diagnostics were obtained with a 39-78-15 architecture (39 neurons in the input layer, 78 in the hidden layer and 15 in the output layer), even the training time is higher (Tab. 2, Tab. 3, Fig.12).



Fig.10. Average square errors and average number of classified diagnostics for different training functions



Fig.11. Average square errors and average number of classified diagnostics for different transfer functions

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Tal	ole	e 2. The a	verag	ge n	umber of	cla	ssified dia	gnostics
from	a	database	with	25	images,	for	different	network
struct	ur	es						

Epochs	Network structure					
$[x10^{3}]$	39-10-15	39-39-15	39-78-15			
1	4,33	15,73	18,27			
2	5,40	17,20	20,67			
3	6,13	19,87	22,53			
4	6,60	20,73	23,27			
5	7,67	21,20	23,53			
6	8,47	21,60	24,00			
7	8,73	22,13	24,67			
8	9,27	22,47	24,80			
9	9,80	22,47	24,93			
10	10,07	22,60	25,00			
11	10,80	22,67	25,00			
12	11,53	22,80	25,00			
13	11,87	22,80	25,00			
14	12,60	22,80	25,00			
15	12,87	22,87	25,00			
16	13,73	23,60	25,00			
17	14,13	23,87	25,00			
18	14,47	23,87	25,00			
19	14,73	24,00	25,00			
20	15,13	24,00	25,00			
21	15,33	24,07	25,00			
22	15,67	24,40	25,00			
23	15,87	24,53	25,00			
24	16,20	24,67	25,00			
25	16,73	24,73	25,00			
26	17,40	24,73	25,00			
27	17,80	24,80	25,00			
28	18,20	24,80	25,00			
29	18,27	24,87	25,00			
30	18,47	24,87	25,00			

The results (square mean errors, number of classified images and training time) are stored in specific files, together with the network configuration and training parameters, so that to be available for later analysis.

Then, more images were added to the database and the training process continued cumulative until all the diagnostics were classified.

If the training continues after this point, a higher precision will be obtained. Unfortunately, this is not useful for our application, because we don't want to recognize a certain image, or a well-known image (as in fingerprint recognition). So, overtraining will decrease the degree of generalization for the network.

Finally, the network was tested with a set of images, some of them representing similar macular diseases or normal macula status. The specificity obtained by the CADx system, regarding the tested database was 90.91%, superior to other applications [13], [14], [31], [32].

Table 3	. The	average	square	mean	error	for	different	
network st	ructur	es						

Epochs	Network structure				
$[x10^{3}]$	39-10-15	39-39-15	39-78-15		
1	0,1493815	0,0588740	0,0448643		
2	0,1366110	0,0437743	0,0290864		
3	0,1255713	0,0324955	0,0198045		
4	0,1155618	0,0255553	0,0138268		
5	0,1073636	0,0211884	0,0105785		
6	0,1001323	0,0180280	0,0079566		
7	0,0941550	0,0155938	0,0051694		
8	0,0883600	0,0133613	0,0037882		
9	0,0829606	0,0121252	0,0027784		
10	0,0776137	0,0110104	0,0020215		
11	0,0715303	0,0100775	0,0014977		
12	0,0670235	0,0093051	0,0011208		
13	0,0634345	0,0086265	0,0008574		
14	0,0601217	0,0080911	0,0006414		
15	0,0562521	0,0076341	0,0004840		
16	0,0534404	0,0063031	0,0003706		
17	0,0511698	0,0053304	0,0002869		
18	0,0492029	0,0047798	0,0002274		
19	0,0473300	0,0042551	0,0001799		
20	0,0457690	0,0039244	0,0001446		
21	0,0437901	0,0036103	0,0001155		
22	0,0418995	0,0030860	0,0000905		
23	0,0400596	0,0025665	0,0000722		
24	0,0378586	0,0019260	0,0000580		
25	0,0359535	0,0016154	0,0000470		
26	0,0340862	0,0014259	0,0000380		
27	0,0325361	0,0011614	0,0000304		
28	0,0312391	0,0010523	0,0000248		
29	0,0302153	0,0008076	0,0000199		
30	0,0292284	0,0007104	0,0000164		



Fig.12. Average square errors and average number of classified diagnostics for different training functions



Fig.13. Image recognition using ANN, from the image processing module

Image recognition can be run from inside the image processing module and inside the diagnosis database module (Fig.13).

The entire image or a certain selected area can be recognized after choosing the proper trained neural network.

A list of four possible diagnostics, ordered by a certitude coefficient, is displayed.

3.5 Patients' database

The doctor receives the decision from the Computer-Aided Diagnosis system and after a discussion with the patient about pathological personal and heredo-collateral antecedents, different medical treatments, establishes the final diagnosis. This can be stored in the patients' database.

Patient's record contains the following information: name of the patient, patient's ID which is used for uniquely identification of the same person, if it is male or female, patient's age, working domain, the date when the record was added or diagnosis date, name and full path of the image file, what the diagnosis is about, a history of patient health status or past diagnosis, the treatment recommended for him and some special remarks [33].

Because of the size of the database, we considered that it is more useful to store only the path and the name of the retina image file. All the image files can be stored in a separate folder on the hard disk for example. It is an advantage because these images can be viewed, analyzed or transferred independently of the database itself, instead of embedding them into the database. From the patients' database, a diagnoses database can be extracted and different types of queries and reports can be generated.

Data have additional value beyond supporting the care of specific patients. For example, subsets of individual patient care data can be used for research purposes, quality assurance purposes, developing and assessing patient care treatment paths (planned sequences of medical services to be implemented after the diagnoses and treatment choices have been made), assessments of treatment strategies across a range of choices, and assessments of medical technologies in use.

The software package is supplied in executable format, so it can be installed on every computer without having to install Matlab®.

4 Conclusion

Entire software for this Computer-Aided Diagnosis system was developed by the authors. The doctor will use the program in a so-called *user* mode, that does not contain all the facilities above, and that can be chosen from the *Settings* menu option. The system is an open one, allowing to create new databases with diagnoses images, new neural networks, to train and optimize them, by an administrator which will have access to all the facilities (*administrator* mode), so that having the possibility to recognize new diagnoses and to continue the training process for improving system performance.

The graphical interface is a user-friendly one.

Starting from classical or digital retina images, image using neural networks recognition algorithms, the Computer-Aided Diagnosis (CADx) system identifies macular diseases with high precision. Images are stored in databases together with patient personal details and treatments and diagnosis information. The software includes image processing modules, databases and neural networks that can be trained for recognizing images of new diseases. The Computer-Aided Diagnosis reduces the doctor's level of incertitude regarding similar diseases, improves the initial and evolutional identification precision of disease. allows monitoring the health status of the patient during new treatment methods, stores images in digital format and generates diagnoses databases that can be used in research, medical practice and specialized teaching.

New development directions for the system will focus on:

- Embedding the system so that to be implemented in a retinal digital investigation camera. This supposes a hardware implementation of the neural network, together with image processing functions;
- On-line retina image analysis on the computer. It will be possible only with a digital retinal investigation camera;
- Integration in a database server for different types of diagnosis, with remote secured access for researchers, doctors, teachers and students. This database will be developed in an important research project called "Researches Concerning the Imaging and Computer-Aided Diagnosis of Human Visual System Diseases", financed by the Romanian Council for Research in Higher Education;
- Adapting the system for using it to recognize images from other domains (industrial for example);
- Improving the system by adding new image features that can be used as inputs in the neural network and testing it on databases containing other images than the retinal ones, in the medical imaging domain;
- The possibility to compare more than two images, for better observing the evolution of a disease, side by side or superposed, using transparency;
- Training neural networks for classification of certain diseases (one network for one type of disease), improving specificity and adapting the CADx software to use multiple networks for recognizing of a large range of diseases.

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