

Fundus characterization for automatic disease screening through retinal image processing

SANDRA MORALES MARTÍNEZ

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Sandra Morales Martínez

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Departamento de Ingeniería Electrónica

Supervisor

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“Nothing in life is to be feared, it is only to be understood. Now is the time to understand more, so that we may fear less.”

Marie Curie

Abstract

The World Health Organization estimates that in 2010 there were 285 million people visually impaired in the world. It is calculated that 80% of these cases are preventable or treatable. In addition, an aging population and an increase in chronic diseases are two factors based on which a higher number of blindness cases is predicted in the future. Hypertension, diabetic retinopathy (DR), age-related macular degeneration (AMD) and glaucoma are the most common pathologies in the current society that provoke retinal damage and can be directly related to blindness and vision loss. The early diagnosis of these diseases allows, through appropriate treatment, to reduce costs generated when they are in advanced states and may become chronic. This fact justifies screening campaigns. However, a screening campaign requires a heavy workload for trained experts in the analysis of anomalous patterns of each disease, which in addition to the increase in population at risk, makes these campaigns economically unfeasible. Therefore, the need for automatic screening system developments is highlighted.

The final goal of this thesis is to develop novel methods that allow the analysis and processing of fundus images to implement an automatic screening of four of the most important diseases affecting world population. In particular, the main objective of the thesis is to build up algorithms for the characterization of the retinal structures and the retina background in order to assist in the discrimination between a “normal” and pathological retina.

Mathematical morphology along with other operators are used for the detection of the retinal vessels and the optic disk. The proposed methods work properly on databases with a large degree of variability. Not only have the main structures been segmented, but significant features have also been extracted to be used in a computer aided diagnosis software for hypertensive risk determination. The texture of the retina background is also analyzed in this work by means of local binary patterns with the aim of identifying DR and AMD and avoiding

the need for segmentation of the characteristic retinal lesions of each disease. The results are promising, above all for AMD diagnosis.

Resumen

La Organización Mundial de la Salud estima que en 2010 había 285 millones de personas con alguna discapacidad visual en el mundo. Se calcula que el 80% de estos casos son evitables o tratables. Además, el envejecimiento de la población y el aumento de las enfermedades crónicas son dos factores que hacen prever un número todavía mayor de casos de ceguera en el futuro. La hipertensión, la retinopatía diabética (RD), la degeneración macular asociada a la edad (DMAE) y el glaucoma son las enfermedades más comunes que provocan daños en la retina y, por tanto, están directamente relacionadas con la ceguera y con la pérdida de visión. El diagnóstico de estas enfermedades en estadios tempranos permite, mediante el tratamiento adecuado, reducir los costes que generan en estados ya avanzados y que en la mayoría de los casos acaban convirtiéndose en crónicas, lo que justifica la realización de campañas de cribado. Sin embargo, una campaña de cribado exige una gran carga de trabajo de personal experto entrenado en el análisis de los patrones anómalos propios de cada enfermedad, lo que sumado al aumento de la población de riesgo, hace que estas campañas sean inviables económicamente. Por lo tanto, se evidencia la necesidad del desarrollo de sistemas de cribado automáticos.

El objetivo final del presente trabajo es la implementación de métodos novedosos de análisis de imágenes de fondo de ojo para usarlos en un sistema de cribado de cuatro de las enfermedades más importantes que afectan a la población actual. En concreto, el objetivo principal de la tesis es el desarrollo de algoritmos para la caracterización de las estructuras y del fondo retiniano, los cuales servirán de ayuda para discriminar una retina “normal” de otra patológica.

Para la detección de los vasos retinianos y del disco óptico, se ha usado morfología matemática además de otros operadores. Se ha demostrado que los métodos propuestos para este fin funcionan adecuadamente en bases de datos con un alto grado de variabilidad. No sólo se han segmentado las principales estructuras retinianas, sino

que, además, se han extraído sus características más significativas para determinar el riesgo hipertensivo. En este trabajo, también se han analizado las texturas presentes en el fondo de la retina por medio de la teoría de los patrones binarios locales con el objetivo de identificar la RD y la DMAE a la vez que se evita la necesidad de la segmentación de las lesiones específicas de cada enfermedad. Los resultados son prometedores, sobre todo, para la detección de la DMAE.

Resum

L'Organització Mundial de la Salut estima que en 2010 havia 285 milions de persones amb alguna discapacitat visual en el món. Es calcula que el 80% d'aquests casos són evitables o tractables. A més, l'envelliment de la població i l'augment de les malalties cròniques són dos factors que fan preveure un número encara major de casos de ceguera en el futur. La hipertensió, la retinopatia diabètica (RD), la degeneració macular associada a l'edat (DMAE) i el glaucoma són les malalties més comuns que provoquen danys en la retina i, per tant, estan directament relacionades amb la ceguera i amb la pèrdua de visió. El diagnòstic d'aquestes malalties en estadis primerencs permet, per mitjà del tractament adequat, reduir els costos que generen en estats ja avançats i que en la majoria dels casos acaben convertint-se en cròniques, la qual cosa justifica la realització de campanyes de garbellament. No obstant això, una campanya de garbellament exigeix una gran càrrega de treball de personal expert entrenat en l'anàlisi dels patrons anòmals propis de cada malaltia, que si es suma a l'augment de la població de risc, fa que aquestes campanyes siguin inviables econòmicament. Per tant, s'evidencia la necessitat del desenrotllament de sistemes de garbellament automàtics.

L'objectiu final del present treball és la implementació de mètodes nous d'anàlisi d'imatges de fons d'ull per a usar-los en un sistema de garbellament de quatre de les malalties més importants que afecten la població actual. En concret, l'objectiu principal de la tesi és el desenvolupament d'algoritmes per a la caracterització de les estructures i del fons retinià, els quals serviran d'ajuda per a discriminar una retina "normal" d'una altra patològica.

Per a la detecció dels vasos retinians i del disc òptic, s'ha usat morfologia matemàtica a més d'altres operadors. S'ha demostrat que els mètodes proposats per a aquest fi funcionen adequadament en bases de dades amb un alt grau de variabilitat. No sols s'han segmentat les principals estructures retinianes, sinó que, a més, s'han extret les seues

característiques més significatives per a determinar el risc hipertensiu. En aquest treball, també s'han analitzat les textures presents en el fons de la retina per mitjà de la teoria dels patrons binaris locals amb l'objectiu d'identificar la RD i la DMAE al mateix temps que s'evita la necessitat de la segmentació de les lesions específiques de cada malaltia. Els resultats són prometedors, sobretot, per a la detecció de la DMAE.

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Chapter 1

Introduction

This chapter presents the motivations behind this thesis, its objectives and its main contributions. In addition, it also introduces the thesis framework and the thesis outline.

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1.1 Motivation

Retinal imaging is nowadays a mainstay of the clinical care and management of patients with retinal as well as systemic diseases. Many important diseases manifest themselves in the retina and originate either in the eye, the brain or the cardiovascular system. In particular, the most prevalent diseases that can be studied via eye imaging and image analysis are: diabetes, diabetic retinopathy, age-related macular degeneration, glaucoma and cardiovascular disease ([Abràmoff et al., 2010](#)).

The World Health Organization (WHO) estimates that in 2010 there were 285 million people visually impaired around the world ([World Health Organization, 2013](#)). In spite of the fact that the number of blindness cases has been significantly reduced in recent years, it is estimated that 80% of the cases of visual impairment are preventable or treatable ([World Health Organization, 2013](#)).

Screening campaigns are usually considered effective if the disease is identified at an early, preferably preclinical, stage and if it is amenable to treatment ([Cree and Jelinek, 2011](#)). The early diagnosis allows, through appropriate treatment, to reduce costs generated when the diseases are in advanced states and may become chronic. In the main, the best approach for the assessment of the population at-risk seems to be a direct, regular and complete ophthalmologic examination ([Herbert et al., 2003](#)). However, a screening campaign requires a large workload of trained experts in the analysis of anomalous patterns of each disease, which in addition to population growth, ageing, physical inactivity and rising levels of obesity make these campaigns economically infeasible because the number of ophthalmologists for the assessment by direct examination of the population at-risk is high ([Verma et al., 2002](#)). Moreover, it must be stressed that this type of retinal disease identification that is based on manual observation, is highly subjective ([Anitha et al., 2011](#)). So, for those reasons, the need for automatic screening systems is highlighted and they have

become increasingly important. Thanks to high resolution of the current retinal cameras, the fundus images can be automatically processed by a screening software providing invaluable help to clinicians in diagnosis and disease prevention. Automatic screening systems are able to relieve physicians of repetitive work, increase efficiency and provide remarkable cost savings in addition to be accessible in rural and remote areas where there is a lack of eye specialists ([American Diabetes Association, 2008](#)).

1.2 Objectives

The final goal of this thesis is the development of new methods that allow the analysis and processing of fundus images for using this information in the implementation of an automatic screening of four of the most important diseases that affect world population: hypertension, diabetic retinopathy, age-related macular degeneration and glaucoma. In particular, the main objective of the thesis is to build up algorithms for the characterization of the retinal structures and the retina background in order to assist in the discrimination between a “normal” and pathological retina. For that, several methods and methodologies have been carried out, which will be further explained. From the main objective, several secondary objectives arise:

- Determination of the characteristic patterns of each disease under study.
- Development and implementation of methods for the location of the main structures of the retina on fundus images which can be related to several pathologies.
- Development and implementation of methods for the extraction of determinant features of the fundus images for diagnosis.
- Selection of a model for the classification of the fundus images as belonging to healthy subject or suspicious of some of the pathologies under study.

1.3 Main contributions

This thesis provides novel methods to characterize fundus images by means of advanced image processing techniques. One of its main contributions is to perform an in-depth study about the use of mathematical morphology in order to detect the most significant retinal structures: vascular network and optic disk. The methods presented in the Chapter 3 are defined by combining different morphological operators, basic ones and other more complex, and by making use of a variant of the watershed transformation, the stochastic watershed, which avoids over and under-segmentation problems related to the classical watersheds.

The vascular network is characterized in two different ways, by means of the segmentation of the complete vasculature and through the vessel centerline extraction. Both approaches are based on mathematical morphology but also use principal curvature information. On the one hand, the main advantage of the method proposed for complete vasculature segmentation is that the optic disk border is not misclassified as vessel which would provoke vessel tracking problems (Morales et al., 2012). On the other hand, the benefit of directly extracting the vessel centerline through the presented algorithm, instead of after a skeletonization process, is that the dependence of previous stages is avoided at the same time that its robustness is kept (Morales et al., 2014a).

The method developed for optic disk segmentation is also based on mathematical morphology although includes a principal component analysis (PCA) in the pre-processing stage to deal with the problem of RGB to gray conversion. It is demonstrated that the use of PCA improves the results of the red component, which is the component most used in the state of the art for the optic disk detection. The extensive validation of this method must also be highlighted. Its results are compared with 5 public databases which shows the high reliability of the algorithm (Morales et al., 2013).

Other contribution of the thesis is that not only are the main retinal structures (blood vessels and optic disk) segmented, but the vascular network is also characterized through the detection of its significant points and the computation of the vessel caliber and the bifurcation angles. These features are used in a computer-aided diagnosis software for hypertensive risk determination which increases the percentage of cases detected if it is compared with the diagnosis obtained from the visual inspection of the fundus (Morales et al., 2014b).

In addition to the hypertension risk determination, the thesis focuses also on the diagnosis of diabetic retinopathy and age-related macular degeneration. The last contribution of the thesis is related to the detection of these diseases. Instead of relying on the segmentation of the characteristic retinal lesions of each disease as most of state-of-the-art works, the diagnosis is based on a texture analysis of the retina background followed by machine learning techniques which achieves promising results.

1.4 Thesis framework

This thesis stands within the framework of two research projects: “Study of retinal microvascular architecture through fundus image processing acquired by non-mydratic retinal cameras” (IMIDTA/2010/47) and “Fundus image processing for automatic screening of ophthalmological diseases - ACRIMA” (TIN2013-46751-R). These projects were or are being funded by the *Instituto de la Mediana y Pequeña Industria Valenciana* (IMPIVA) and by the *Ministerio de Economía y Competitividad* of Spain, respectively.

The first one was carried out thanks to the collaboration among LabHuman (*Univesitat Politècnica de València*), the Department of Pediatrics of the *Hospital General de Valencia* and the *Fundación Oftalmológica del Mediterráneo*. The aim of this project was the study of retinal microvascular architecture by measuring the bifurcation angles and the caliber of the retinal arterioles in children with low

birth weight, as a prognostic marker of hypertension and cardiovascular pathology. For that purpose, a computer-aided diagnosis software, able to segment blood vessels and perform angle and caliber measurements, was developed. This project supposed the beginning of a research line in LabHuman about retinal image processing.

The scope of ACRIMA is wider than the previous project. Its objective is to develop an automatic screening system for the three most significant diseases related to visual impairment: glaucoma, diabetic retinopathy and age-related macular degeneration. At present, there is no screening system as the proposed in the project, which combines the screening of these three diseases. The members of the *Fundación Oftalmológica del Mediterráneo*, renamed nowadays as *FISABIO-Oftalmología Médica*, continue collaborating with LabHuman in this project.

Both projects have a common denominator. The resulting systems are designed to be used by general practitioners (GP) as computer-aided diagnosis tools. When a patient goes to a medical consultation, the GP will determine the need for medical tests. If they are needed, a fundus image of the patient will be captured and processed by the developed tools. If the tool detects that there is pathology risk, the patient will be sent to the ophthalmologist who will decide if some specific tests are needed and will determine the final diagnosis. This process is summarized in Figure 1.1.

1.5 Outline

This thesis is divided in 5 chapters. This chapter has presented the motivations behind the research involved in the thesis, the main and secondary objectives and its main contributions. Finally, the thesis framework is also explained.

Chapter 2 introduces the anatomical aspects of the retina, the most important retinal diseases and the most relevant groups working on the

development of screening systems for the detection of some of these diseases.

Chapter 3 presents new methods to detect the retinal vessels and the optic disk, two of the most important anatomical structures of the fundus, and to determine some of their main characteristics. These elements will be key for pathology detection.

In Chapter 4, the texture of the retina background is analyzed by means of local binary patterns to differentiate between pathological and healthy images. With this analysis, it is intended to determine if a patient is suspicious of pathology but avoiding the need for segmentation of different types of lesions.

General and final conclusions in addition to future prospects are presented in Chapter 5. Next, the scientific publications derived from this thesis are cited.

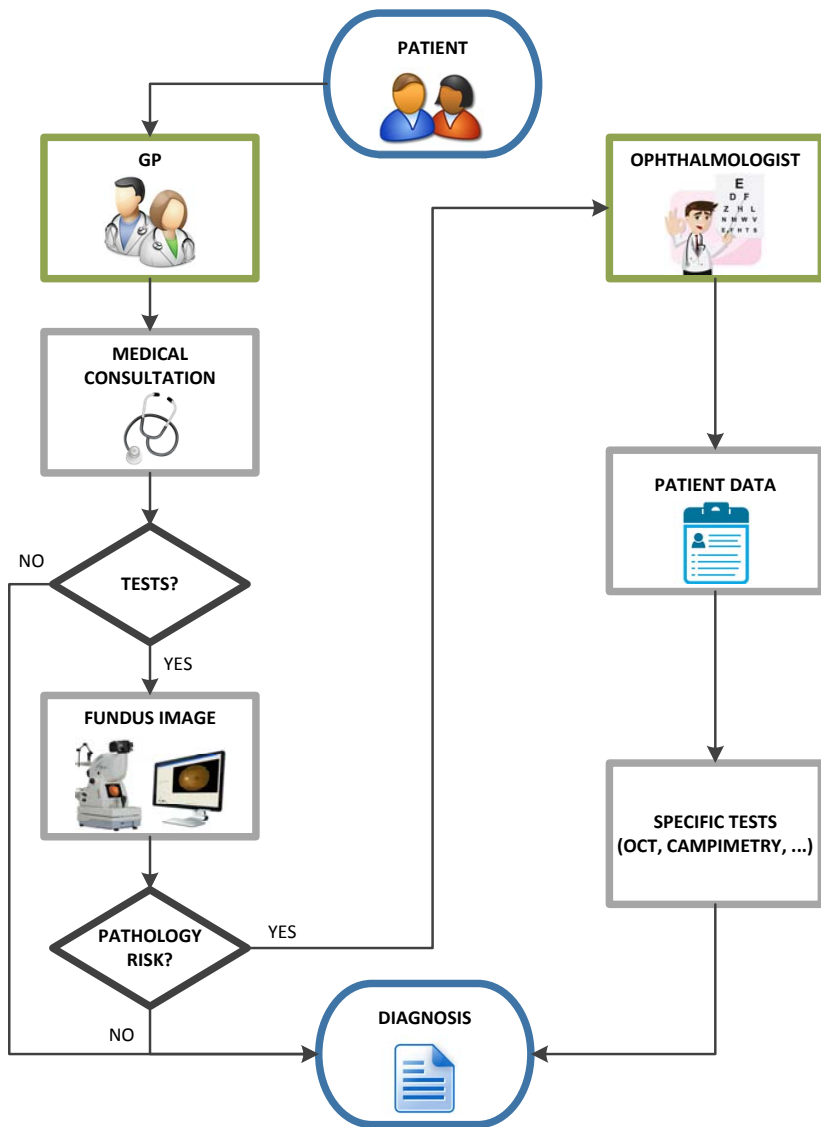


Figure 1.1: Thesis framework flowchart.

Chapter 2

The retina

This chapter presents a brief explanation about anatomical aspects of the retina, the most important retinal diseases and the most relevant groups working on the development of screening systems for the detection of some of these diseases.

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2.1 Retina's anatomy

The eye is the sense organ of sight. It is often compared to a camera which captures and transmits light through a lens to create an image.

The eyeball consists of three coats: the sclera, with its anterior part, the cornea; the uvea that includes the choroid, ciliary body and iris; and the retina (Figure 2.1). The retina is the innermost layer and the focus of this thesis. The purpose of the retina is to transform light into a nerve impulse.

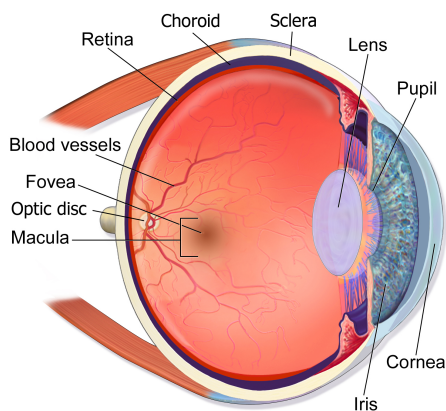


Figure 2.1: *Eyeball anatomy.*¹

Approximately 1.2 million axons of ganglion cells converge to form the optic nerve. The optic nerve is directly connected to the retina. Its function is to send the nerve impulses produced by the retina to the brain to be interpreted and transformed into a visual perception.

The retina is made up two kinds of photoreceptors (cones and rods). The cones are responsible for bright light and colors and the rods for peripheral and night vision. There are 5 million cones and 125 million rods in an average retina.

¹Image adapted from Blausen gallery 2014 - Wikiversity Journal of Medicine (http://en.wikiversity.org/wiki/Blausen_gallery_2014).

Next, the most important structures of the retina will be detailed. They can be observed in Figure 2.2.

- Blood vessels.

Blood vessels supply the retina. They enter and exit to and from the retina via the optic disk. The retinal vascular network is composed by arteries and veins. The main branches of the central retinal artery and vein form four vascular arcades.

- Optic disk.

The optic disk is a circular to oval area where the optic nerve is connected to the retina. It measures about 2 x 1.5 mm across and can be seen as a bright yellow disk. It is also called blind spot because there are not receptors in this part of the retina.

In the center of the optic disk there is a white depression with cup-like shape called optic cup. The optic cup is not occupied by nerve fibers, i.e., devoid of neuroretinal tissue. The neuroretinal rim is the area the optic disk which contains the neural elements and is located between the edge of the disk and the optic cup. The cup-to-disk ratio is important clinically because its increase indicates the death of ganglion cells.

- Macula.

The macula is a pigmented area near the center of the retina that is responsible for finest central vision. It is about 1.5 mm in diameter and it is located two disk diameter temporal to optic disk. The normal macula appears darker than the rest of the retina.

- Fovea.

The fovea is a small pit (approximately 500 μm in diameter) involved with high acuity vision. It is located in the center of the macula and contains a high density of cones. Rods are present everywhere in the retina except at the fovea.

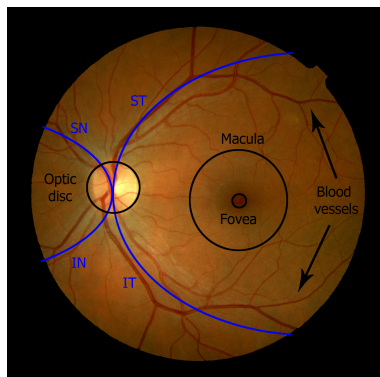


Figure 2.2: *Main parts of the retina: blood vessels, optic disk, macula and fovea. The vascular arcades formed by the blood vessels are marked in blue: superotemporal (ST), inferotemporal (IT), superonasal (SN) and inferonasal (IN).*

2.2 Pathologies related to the retina

Due to the retina's vital role in vision, the damage of the retina can cause visual impairment and, even, permanent blindness. Hypertension, diabetic retinopathy, age-related macular degeneration and glaucoma are some of the most common diseases in the current society that provoke retinal damage and are detectable through fundus inspection ([World Health Organization, 2010a](#); [Abràmoff et al., 2010](#)). Next, these pathologies will be described, emphasizing the characteristic patterns of the disease and summarizing the works found in the literature for their detection. By way of introduction, in [Figure 2.3](#) the effects of some diseases on the vision are shown.

2.2.1 Hypertensive retinopathy

Hypertension is a chronic disease characterized by a continuous increase in the blood pressure in the arteries. It is one of the major cardiovascular risk factors. Globally, the overall prevalence of high blood pressure in adults aged 25 and above was around 40% in 2008

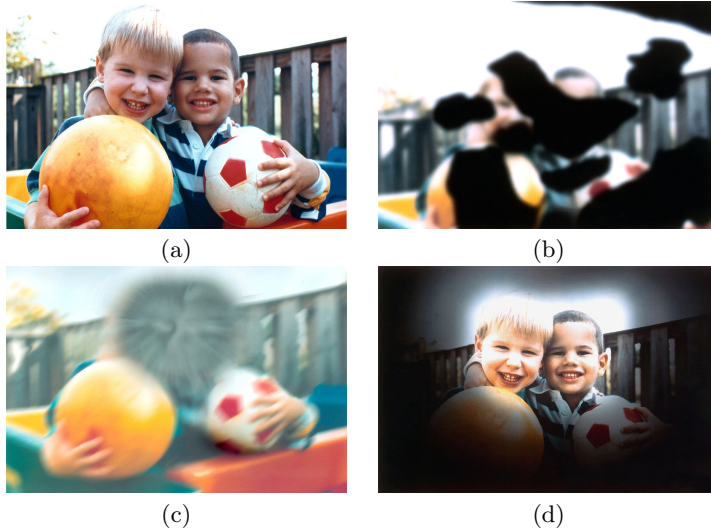


Figure 2.3: *Effects of different pathologies on the vision: (a) Normal vision, (b) With diabetic retinopathy, (c) With age-related macular degeneration and (d) With glaucoma.*²

([World Health Organization, 2010b](#)). This data is directly related to hypertension prevalence.

Hypertension results from a complex interaction of genes and environmental factors as overweight and salt and alcohol intake. Moreover, as blood pressure rises with aging, the risk of becoming hypertensive increases in later life ([Vasan et al., 2002](#)).

Although hypertension is rarely accompanied by visual symptoms, it can be suspected on the basis of the presence of hypertensive retinopathy since changes in the arterial hypertension of the retina are indicator of the state of the blood vessels in other organs ([Wong and Mitchell, 2007](#)). In particular, vascular changes produced in

²Images provided by the National Eye Institute (www.nei.nih.gov).

systemic diseases usually induce specific modifications in the retinal vessels, such as: generalized retinal-arteriolar narrowing which implies a decrease in the ratio between the diameter of retinal arteries and veins, arteriovenous nicking, focal arteriolar narrowing, alteration in the arteriolar light reflex, arteriolar tortuosity, an increase in the angle of arteriolar branching, central or branch retinal artery and vein occlusions, etc. (Wong et al., 2001; Liew and Wang, 2011). Some of these signs can be observed in Figure 2.4.

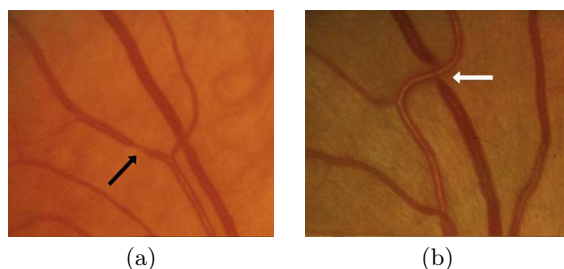


Figure 2.4: *Some retinal vascular signs: (a) Focal arteriolar narrowing and (b) Arteriovenous nicking. Images extracted from (Liew and Wang, 2011).*

In general, state-of-the-art software applications that compute measurements of retinal vessels for hypertensive risk determination are focused only on caliber determination and no other measures are performed. At the beginning, software applications were not used to detect retinal vessels or measure geometrical properties automatically but the diameters of all arterioles and venules were measured by trained graders using different tools provided by the systems to make this task simpler (Wong et al., 2004; Knudtson et al., 2003; Hubbard et al., 1999). Nowadays, there are some automatic or semiautomatic approaches to perform these measures (Ruggeri et al., 2007; Muramatsu et al., 2010; Tramontan et al., 2008; Niemeijer et al., 2011; Ortega et al., 2010).

2.2.2 Diabetic retinopathy

Diabetes is a metabolic disease due to insufficient production of the hormone insulin, or a resistance to its action which prevents the glucose uptake of the blood by the cells of the organism giving place to high blood sugar levels. Diabetic retinopathy (DR) is a common symptom of diabetes and one of the leading causes of blindness worldwide. In fact, this disease occurs in 80% of patients who have had diabetes for more than 10 years (Wang et al., 2000). To ensure that DR treatment is received on time, fundus images from diabetic patients must be examined at least once a year (Fong et al., 2003).

Diabetic retinopathy is a silent disease because only affects vision when retina changes have progressed to a level where treatment is complicated and expensive and the maintenance of vision is uncertain. The most effective treatment of the disease can only be administered in the early stages and, thus, early detection through regular screening is of paramount importance. For automatic screening of DR, most systems are based on the detection of the following features (Yun et al., 2008):

- **Microaneurysms.** Microaneurysms are the first identifiable characteristic of the disease. They are small saccular dilatations of capillaries that appear as round spots of dark red color.
- **Exudates.** Exudates are accumulations of lipids and proteins in the retina. They are typically bright lesions of white or cream color.
- **Hemorrhages.** Hemorrhages become visible when the DR degree progresses. They indicate an ischemia increase (lack of oxygen) of the retina.

DR can be broadly classified as nonproliferative DR (NPDR) and proliferative DR (PDR) depending on the presence and amount of the mentioned features (Early Treatment DR Study Research Group, 1991).

Figure 2.5 identifies the different types of DR features and Figure 2.6 shows two fundus images with NPDR and PDR, respectively.

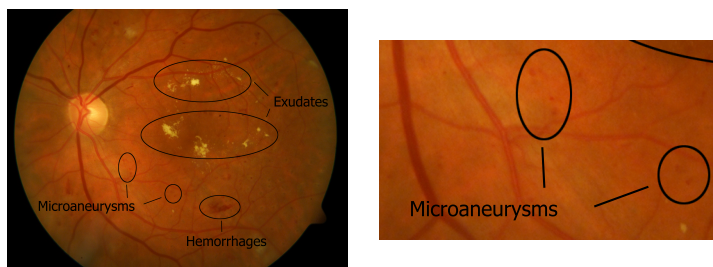


Figure 2.5: *Features of diabetic retinopathy (zoom in the area with microaneurysms).*

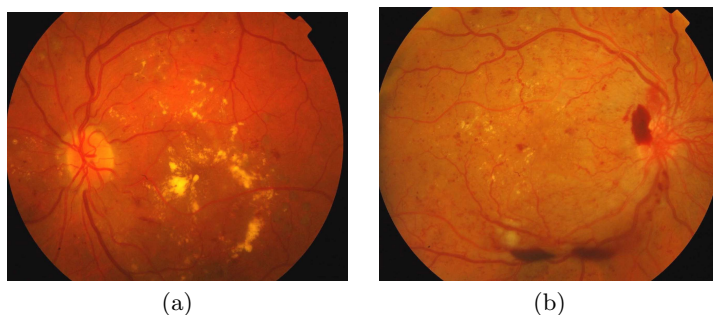


Figure 2.6: *Grades of diabetic retinopathy: (a) Nonproliferative DR (also known as background DR) and (b) Proliferative DR.*

DR is the disease which more research groups are working on. So, DR and normal retina have been classified for multitude of methods with different results up to now. Some of them will be enumerated. Multilayer perceptron neural network is used in (Sinthanayothin et al., 2002) achieving a sensitivity of 80.21% and a specificity of 70.66%. The identification of exudates, hemorrhages and microaneurysms is needed to carry out the screening in (Singalavaniya et al., 2006), which gets a sensitivity of 74.8% and a specificity of 82.7%. Microaneurysms

are detected by Bayes optimality criteria and early detection of DR is performed by a decision support system in (Kahai et al., 2006) obtaining a sensibility of 100% but a specificity of 67%. An automated grading for DR detection with 90.5% sensitivity and 67.4% specificity is used in (Philip et al., 2007). The classification is based on fovea, blood vessels, optic disk and bright and dark lesion segmentations in (Estabridis and de Figueiredo, 2007) obtaining 90% of accuracy. A sensitivity of 84% and a specificity of 64% was achieved by the system proposed in (Abràmoff et al., 2008). Note that these works are validated on different databases therefore their direct comparison is difficult. A wider review about DR detection systems can be found in (Faust et al., 2012).

2.2.3 Age-related macular degeneration

Age-related macular degeneration (AMD) is a degenerative condition which degrades progressively the macula. The proper functioning of the macula lets appreciate details, reading, driving, face recognizing, etc. AMD is a common eye disease that affects adults and it has become the leading cause of blindness over 60 years (Jager et al., 2008). Its progression is so slow that patients only notice small changes in their vision although it can provoke a total loss of central vision. In spite of the fact that AMD has currently no definitive cure, early detection allows to provide treatments that retard and mitigate the effects of the disease (Jager et al., 2008).

Drusen, yellow deposits in the macula, are one of the early signs more common of the AMD and the basis on which ophthalmologists pay attention to detect the disease. Drusen usually are of two types: hard and soft. Hard drusen are small, round, with sharp edges. These drusen are generally not harmful if they present in small quantity. Soft drusen are larger than the hard drusen and may have indistinct borders. As the disease progresses, the number and size of the drusen increases and other signs can also appear such as geographic atrophy of the retinal pigment epithelium, choroidal neovascularization, disciform scar, etc.

(Miller, 2013). In Figure 2.7, different signs of AMD can be observed.

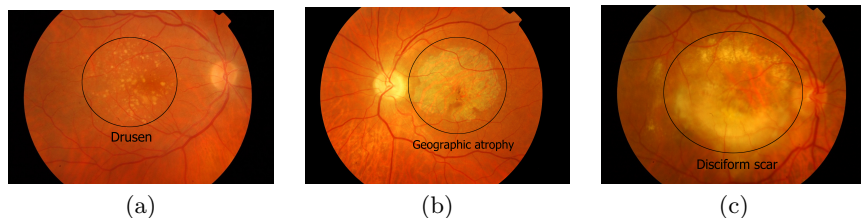


Figure 2.7: Signs of age-related macular degeneration: (a) Drusen, (b) Geographic atrophy of the retinal pigment epithelium and (c) Disciform scar.

Most works of the state of the art related to AMD is based on drusen segmentation but not in AMD diagnosis (Hijazi *et al.*, 2012) although some reports can be found (Agurto *et al.*, 2011a; Garnier *et al.*, 2014). Agurto *et al.* use an Amplitude Modulation - Frequency Modulation (AM-FM) technique to define the features followed by two-step Partial Least Squares (PLS) classifier achieving a sensitivity of 94% and a specificity of 50%. Garnier *et al.* do not perform drusen segmentation but they analyze texture information through wavelet decomposition and local binary patterns. A sensitivity of 91% and a specificity of 95% are obtained using a LDA classifier. These methods are validated on different private databases.

2.2.4 Glaucoma

Glaucoma is a progressive disease of the optic nerve caused by high intraocular pressure due to a bad drainage of the ocular fluid. Clinically, it causes a progressive and irreversible loss of the visual field that progresses to the complete loss of vision. It is currently the second leading cause of blindness worldwide and affects one in every hundred people under 50 and one in ten over 80 years (Kwon *et al.*, 2009). In most cases, glaucoma is belatedly detected when visual field loss is irreversible. Currently, there is no cure for glaucoma damage and,

therefore, early detection and prevention is the only way to prevent progression to total vision loss.

According to the literature, the diagnosis of glaucoma is mainly based on three studies: the measurement of intraocular pressure by tonometry, the examination of the visual field by campimetry and the computing of the ratio between the optic cup and optic disk area from fundus images, also known as cup-to-disk ratio (CDR) (Nath and Dandapat, 2012). Since this thesis focuses on the detection of glaucoma by means of retinal image analysis, it is the last of the studies, the measure of the cup-disk ratio, which is of our interest. The optic nerve damage is manifested in a change in the appearance of the optic cup, so that its size (and therefore the CDR) is related to the presence or absence of glaucoma. For healthy patients, CDR falls in the range of 0.3 to 0.5 and for glaucoma it is greater than 0.5. The sight of vision is completely lost at the CDR value of 0.8 (Nath and Dandapat, 2012). In Figure 2.8 the existing difference between a fundus with and without glaucoma can be appreciated.

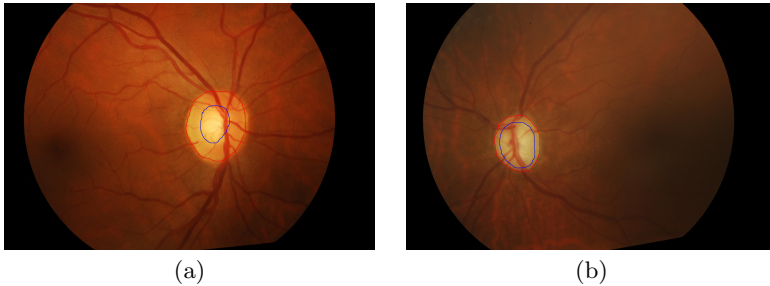


Figure 2.8: *Glaucoma effects on the cup-to-disk ratio: (a) Healthy fundus and (b) Glaucomatous fundus. Optic disk is marked in red and optic cup in blue.*

Among the different works that deal with glaucoma detection the following can be stood out. The ARGALI system performs an automatic cup-to-disk ratio measurement using level-sets and achieves a RMS error

of 0.05 with a risk assessment accuracy of 95% (Liu et al., 2009). Nayak *et al.* not only use the CDR as feature but also take into account other anatomical ratios (the ratio of the distance between optic disk center and optic nerve head to diameter of the optic disk and the ratio of blood vessels area in inferior-superior side to area of blood vessels in the nasal-temporal side). All these features are incorporated in a neural network classifier which obtains a sensitivity and specificity of 100% and 80% respectively (Nayak et al., 2009). Hatanaka *et al.* carried out a different approach and measure the cup-to-disk ratio through the vertical profile of the optic disk getting a sensitivity of 80% and a specificity of 85% (Hatanaka et al., 2010). These statistics were computed on different databases. More techniques of glaucoma detection can be found in (Nath and Dandapat, 2012).

2.3 Relevant groups

The most relevant groups working on retinal image processing, which have developed some prototype of detection systems of one or more pathologies under study in this thesis, are:

International groups

The Retinal Image Computing & Understanding group³ at the University of Lincoln (United Kingdom) works on segmentation and measurement of the retinal vessels (Lowell et al., 2004; Al-Diri et al., 2009; Al-Diri and Hunter, 2009) and optic disk segmentation (Lowell et al., 2004). Their works are mainly related to hypertension although some publications are referred to diabetic retinopathy (Massey et al., 2009). Nowadays, they are the coordinators of a European project, named “REVAMMAD - REtinal Vascular Modelling, Measurement and Diagnosis”⁴, included in the Seventh Framework Programme.

³www.aldiri.info/Default.aspx

⁴<http://revammad.blogs.lincoln.ac.uk/revammad/>

VisionQuest Biomedical⁵ (Albuquerque) focuses on diabetic retinopathy screening although they also work on the detection of other pathologies such as age-related macular degeneration (Agurto et al., 2011a; Agurto et al., 2011b). Their system EYESTAR⁶ is now in a clinical trial.

The Institute for Infocomm Research⁷ (Singapore). They have mainly worked on glaucoma detection through a system for automatic measurement of the cup-to-disk ratio called ARGALI (Liu et al., 2009). They have also worked, along with the National University of Singapore, on the development of a platform for vascular disease risk prediction (Hsu et al., 2011).

The Computer Vision and Image Processing (CVIP) group⁸ at the University of Dundee (Scotland) has worked on the implementation of VAMPIRE⁹, a vessel assessment and measurement platform (Perez-Rovira et al., 2011).

The University of Aberdeen (Scotland) has worked on an automated grading of diabetic retinopathy in primary care screening (Philip et al., 2007; Goatman et al., 2011).¹⁰ Although only for a disease, it is the only existing CE-accredited commercial system in terms of analysis for automatic fundus screening.¹¹

⁵www.visionquest-bio.com

⁶www.visionquest-bio.com/eyestar-tm.html

⁷www.i2r.a-star.edu.sg/industries/medical-and-healthcare

⁸<http://cvip.computing.dundee.ac.uk/>

⁹<http://vampire.computing.dundee.ac.uk/>

¹⁰www.abdn.ac.uk/heru/research/assessment-of-technologies/projects/the-role-of-automated-grading-of-diabetic-retinopa/

¹¹www.shil.co.uk/Products/igradingtm-platform-diabetic-retinopathy-screening-software.html

National groups

The VARPA group¹² at University of A Coruña works mainly on hypertension detection by means of automatic computation of the arteriolar-to-venular ratio. They developed a web application for this purpose known as SIRIUS (Ortega et al., 2010).

The University of Huelva works on an expert system for early automated detection of diabetic retinopathy.¹³ At this moment, there are not results of this system but some algorithms for retinal structure segmentation have been published (Aquino et al., 2010; Marín et al., 2011).

In spite of the amount of groups working on retinal image processing, there is no system that detects at the same time the most common diseases that cause retinal damage. Moreover, the screening systems developed up to now are characterized by achieving high sensitivity (correct detection of pathological retinas) but not specificity enough (correct detection of healthy subjects) to be effective for screening purposes. All these facts stand out the need to continue working on this topic.

¹²www.varpa.es

¹³www.uhu.es/retinopathy/index.php

Chapter 3

Retinal structure characterization

The retinal vessels and optic disk are two of the most important anatomical structures of the fundus. This chapter presents new methods to detect them and to determine some of their main characteristics. The detection and analysis of these elements are used for hypertensive risk determination. Finally, results and conclusions of the chapter are detailed.

In this chapter, works previously published, although adapted for this thesis, are included:

- *Subsection 3.3.1 - Vessel extraction.*

Morales, S., Naranjo, V., Angulo, J., Fuertes, J. J., Alcañiz, M. (2012). Segmentation and Analysis of Retinal Vascular Tree from Fundus Images Processing. In *International Conference on Bio-inspired Systems and Signal Processing (BIOSIGNALS 2012)*, pages 321-324.

- *Subsection 3.3.2 - Centerline extraction.*

Morales, S., Naranjo, V., Angulo, J., López-Mir, F., Alcañiz, M. (2014). Determination of Retinal Network Skeleton through Mathematical Morphology. In *22nd European Signal Processing Conference (EUSIPCO 2014)*, pages 1691-1695.

- *Section 3.4 - Significant points.*

Morales, S., Naranjo, V., Colomer, A., Alcañiz, M. (2015). Significant point characterization in fundus images. In *5th International Conference on Image Processing Theory, Tools and Applications (IPTA 2015)* (Accepted for publication).

- *Section 3.5 - Optic disk segmentation.*

Morales, S., Naranjo, V., Angulo, J., Alcañiz, M. (2013). Automatic detection of optic disc based on PCA and mathematical morphology. *IEEE Transactions on Medical Imaging*, 32(4): 786-796.

- *Section 3.6 - Application to pathology detection: Hypertensive risk.*

Morales, S., Naranjo, V., Navea, A., Alcañiz, M. (2014). Computer-Aided Diagnosis Software for Hypertensive Risk Determination Through Fundus Image Processing. *IEEE Journal of Biomedical and Health Informatics*, 18(6):1757-1763.

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3.1 Introduction

Retinal structure segmentation is a fundamental component of most automatic retinal disease screening systems (Fraz et al., 2012). It is usually a prerequisite before systems carrying out more complex tasks as identifying different pathologies. In general, several anatomical structures are segmented through fundus image processing and then some features are extracted from them to characterize each pathology. The most important anatomical structures of the fundus are the vascular network that is the retinal blood vessels, and the optic disk (OD). Both of them can be used as reference to detect other anatomical ocular structures or several retinal lesions and to identify some fundus features. For example, among their multiples uses, the following can be stressed. Morphological attributes of retinal blood vessels, such as length, width, tortuosity and/or branching pattern and angles are utilized for diagnosis, screening, treatment, and evaluation of various cardiovascular and ophthalmologic diseases (Kanski and Bowling, 2011). The relation between the size of the optic disk and the cup (cup-to-disk ratio) is widely used for glaucoma diagnosis (Hatanaka et al., 2010). The relatively constant distance between the optic disk and the fovea is useful for estimating the location of the macula, area of the retina related to fine vision (Niemeijer et al., 2009). Their detection may also be used for not taking them into account in the analysis of the retina background with the aim of pathology diagnosis, as discussed in Chapter 4. Due to their importance in screening systems of ophthalmologic diseases, this chapter presents new algorithms related to detection and characterization of the retinal vessels and the optic disk. Their performance and their place with respect to previous studies will be detailed along the chapter.

Numerous segmentation methods of retinal structures have been reported in the literature. Nevertheless, only a review of the state of the art of the methods regarding the description of the vascular tree and the optic disk will be performed here.

Referring to vessel extraction techniques, they can be mainly divided into four categories: edge detectors, matched filters, pattern recognition techniques and morphological approaches. A more extensive classification can be found in (Fraz *et al.*, 2012). Most edge detection algorithms assess changes between pixels values by calculating image gradient magnitude and then it is thresholded to create a binary edge image (Martinez-Perez *et al.*, 2007; Jiang and Mojon, 2003). Matched filters are filters rotated in different directions in order to identify the cross section of blood vessels (Hoover *et al.*, 2000; Al-Rawi *et al.*, 2007). Pattern recognition techniques can be divided into supervised and unsupervised approaches. Supervised methods, such as artificial neural networks (Sinthanayothin *et al.*, 1999) or support vector machines (Ricci and Perfetti, 2007; Marín *et al.*, 2011), exploit some prior labelling information to decide whether a pixel belongs to a vessel or not, while unsupervised algorithms perform the vessel segmentation without any prior labelling knowledge (Kande *et al.*, 2010). Morphological processing is based on vessels characteristics known a priori (line connected segments) and combines morphological operators to achieve the segmentation (Zana and Klein, 2001; Mendonça and Campilho, 2006; Sun *et al.*, 2011). Although most of the state-of-the-art methods look for detecting all vessel pixels of the vascular tree, there are also some attempts based on finding the vessel skeleton or, in other words, the vessel centerline. The work of Chen *et al.* is based on shortest path connection (Chen *et al.*, 2011), Sofka and Stewart on the use of matched filters (Sofka and Stewart, 2006), Wu and Derwent on ridge descriptors (Wu *et al.*, 2009) and Walter and Klein and Bessaid *et al.* on the application of watershed transformation (Walter and Klein, 2002; Bessaid *et al.*, 2009).

Once the vessels are segmented, the identification of the vascular bifurcations and crossovers is helpful for predicting cardiovascular diseases and can be used as biometric features and for image registration (Grisan *et al.*, 2004; Bevilacqua *et al.*, 2005; Bhuiyan *et al.*, 2007; Calvo *et al.*, 2011; Azzopardi and Petkov, 2011). After identifying the vascular bifurcations, the angles generated in these points are usually measured

by fitting the vessels that form the bifurcation by straight lines and then estimating the angle between them (Gao et al., 2000; Martinez-Perez et al., 2002; MacGillivray et al., 2012).

Regarding the OD-boundary detectors, the presented techniques can mainly be grouped into template-based methods, deformable models, and morphological algorithms. Different approaches have been proposed according to template-based methods: edge detection techniques followed by the Circular Hough Transform (Park et al., 2006; Aquino et al., 2010); pyramidal decomposition and Hausdorff-based template matching (Lalonde et al., 2001); color decorrelated templates (Kauppi and Kälviäinen, 2008); or a kNN-regressor along with a circular template (Niemeijer et al., 2009). Concerning deformable models, GVF-snake (Osareh et al., 2002), ASM (Li and Chutatape, 2004) and modified active contours, which exploit specific features of the optic disk anatomy (Lowell et al., 2004) or incorporate knowledge-based clustering and updating (Xu et al., 2007), have also been used to this purpose. As for algorithms based on mathematical morphology, most of them detect the OD by means of watershed transformation, generally through marker-controlled watershed (Walter et al., 2002; Eswaran et al., 2008; Welfer et al., 2010), although each author chooses different markers. The centroid of the largest and brightest object of the image is considered as an approximation for the locus of the OD and it is used as internal marker (Walter et al., 2002). The extended minima transformation (Gonzalez et al., 2004) is applied to select the internal markers and external markers are calculated as an effectively partition of the image into regions, so that each region contains single internal marker and part of the background (Eswaran et al., 2008). A list of pixels belonging to the main vessels arcade in the vicinity of an internal OD point previously detected are also used (Welfer et al., 2010). On the other hand, some authors propose combining various algorithms to get a better approximation of the OD-boundary (Hajer et al., 2008; Qureshi et al., 2012).

The rest of the chapter is organized as follows: Section 3.2

describes the theoretical background. Sections 3.3, 3.4 and 3.5 present the algorithms proposed for detection and characterization of the main retinal structures. Section 3.6 shows how these methods can be applied for hypertensive risk determination. Section 3.7 details the experimental results of the methods explained above and their comparison with other state-of-the-art algorithms. Finally, Section 3.8 provides conclusions.

3.2 Theoretical background

This section focuses on introducing the main concepts that will be used in the chapter in order to facilitate its comprehension to the reader. The main morphological operators and the stochastic watershed transformation will be described below to be able to understand the methods developed. More complete readings about mathematical morphology applied to digital image processing can be found in (Serra, 1982; Soille, 2003).

3.2.1 Morphological operators

Mathematical morphology is a non-linear image processing methodology based on minimum and maximum operations whose aim is to filter or extract relevant structures of an image (Serra, 1982). Next, different morphological filters will be defined and an explanation about geodesic transformations will also be included. The different operators will be illustrated with figures along the section.

Morphological filters

Let f be a gray-scale image which is defined as $f(\mathbf{x}) : E \rightarrow T$ where \mathbf{x} is the pixel position. In the case of discrete-valued images, $T = \{t_{min}, t_{min} + 1, \dots, t_{max}\}$ is an ordered set of gray-levels. Typically, in digital 8-bit images $t_{min} = 0$ and $t_{max} = 255$. Furthermore, let $B(\mathbf{x})$

be a sub-set of Z^2 called structuring element (SE) centered at point \mathbf{x} , whose shape is usually chosen according to some a priori knowledge about the geometry of the relevant and irrelevant image structures. The two basic morphological operators are:

$$\begin{aligned} \text{Dilation: } [\delta_B(f)](\mathbf{x}) &= \max_{b \in B(\mathbf{x})} f(\mathbf{x} + \mathbf{b}) \\ \text{Erosion: } [\varepsilon_B(f)](\mathbf{x}) &= \min_{b \in B(\mathbf{x})} f(\mathbf{x} + \mathbf{b}). \end{aligned} \quad (3.1)$$

Their purpose is to expand light or dark regions, respectively, according to the size and shape of the structuring element. See Figure 3.1.

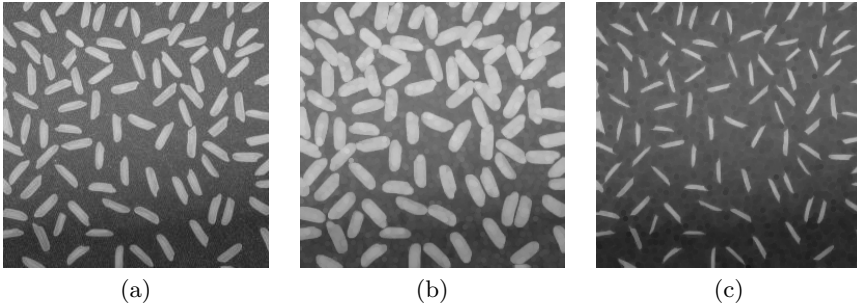


Figure 3.1: Basic morphological operators: (a) Original image, (b) Dilation and (c) Erosion. Both operators were performed using a disk-shaped structuring element of radius 2.

In mathematical morphology, the gradient $\varrho(f)(\mathbf{x})$ of an image $f(\mathbf{x})$ can be obtained as the pointwise difference between a unitary dilation and a unitary erosion, i.e.,

$$\varrho(f)(\mathbf{x}) = \delta_B(f)(\mathbf{x}) - \varepsilon_B(f)(\mathbf{x}). \quad (3.2)$$

The goal of the gradient transformation is to highlight image contours.

Dilation and erosion can be combined successively to obtain a new set of operators or basic filters given by:

$$\begin{aligned} \text{Opening: } \gamma_B(f) &= \delta_B(\varepsilon_B(f)) \\ \text{Closing: } \varphi_B(f) &= \varepsilon_B(\delta_B(f)). \end{aligned} \tag{3.3}$$

Light or dark structures are respectively filtered out from the image by these operators regarding the structuring element chosen (Figure 3.2).

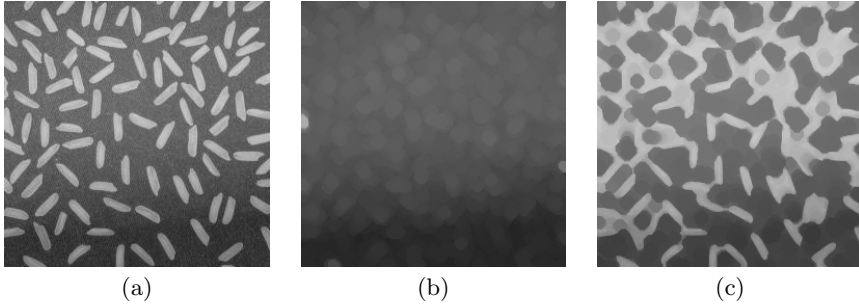


Figure 3.2: Basic morphological filters: (a) Original image, (b) Opening and (c) Closing. Both filters were performed using a disk-shaped structuring element of radius 5.

Openings and closings act as non-linear low-pass filters simplifying the details of an image. However, the objects of interest may also be those details and, not always, only one operator is enough to obtain all the desired objects. To solve these problems there exist other morphological operators derived from the previous ones: top-hats and the supremum of openings or the infimum of closings.

The top-hat transformation aims to suppress slow trends, therefore to enhance the contrast of some features in images, according to size or shape criteria. The top-hat (or positive top-hat) is the residue between the original image and its opening and the dual top-hat (or negative top-hat) can be defined as the residue between a closing and its original

image, i.e.,

$$\begin{aligned} \text{Top-hat: } \rho_B^+(f) &= f - \gamma_B(f) \\ \text{Dual top-hat: } \rho_B^-(f) &= \varphi_B(f) - f. \end{aligned} \tag{3.4}$$

The top-hat transformation is used to extract contrasted components with respect to the background (Figure 3.3(c)). The basic top-hat extracts positive components and the dual top-hat the negative ones.

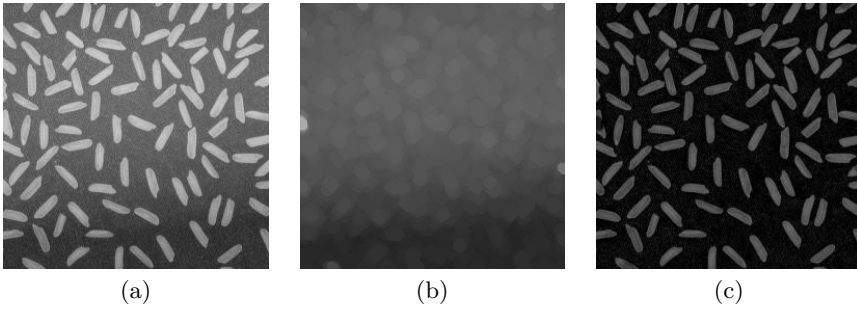


Figure 3.3: Top-hat transformation: (a) Original image, (b) Opening with a circular SE of radius 5 and (c) Top-hat.

For a collection γ_i of openings, the supremum of openings, $\gamma^{sup} = \bigvee_i \gamma_i$ can be considered as the union of all openings which is also an opening. Figure 3.4 shows an example where the supremum of two openings performed with linear SE of different orientations was performed. This is useful when the object to be extracted has not a unique orientation, as in the case of the figure. On the other hand, if φ_i is a collection of closings, then the infimum of closings, $\varphi^{inf} = \bigwedge_i \varphi_i$ is the intersection of all closings which is a closing as well.

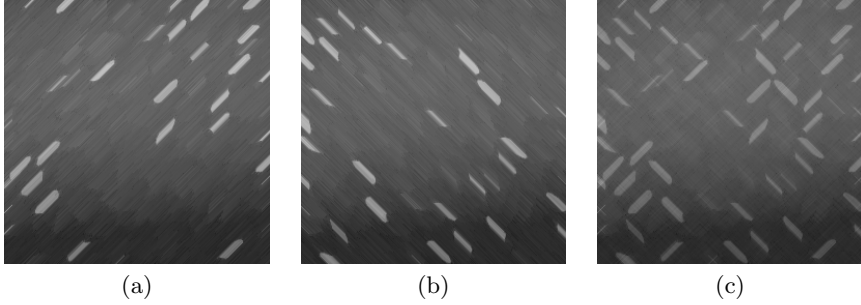


Figure 3.4: *Supremum of openings: (a) Opening with a linear SE (+45°), (b) Opening with a linear SE (-45°) and (c) Supremum of linear openings. The original image is that shown in Figure 3.3(a)*

Geodesic operators

Other operators used in this thesis are: reconstruction by dilation and close-holes. First, elementary geodesic transformations will be introduced to be able to explain these operators. The geodesic transformations involve two images: a marker image f and a reference image g . By definition, the reference image must be greater than or equal to the marker image ($f \leq g$).

The *geodesic dilation* is the iterative unitary dilation of the marker image f with respect to the reference image g ,

$$\delta_g^{(n)}(f) = \delta_g^{(1)}\delta_g^{(n-1)}(f), \text{ being } \delta_g^{(1)}(f) = \delta_B(f) \wedge g. \quad (3.5)$$

The *reconstruction by dilation* is the successive geodesic dilation of f regarding g up to idempotence,

$$R_g^\delta(f) = \delta_g^{(i)}(f), \text{ so that } \delta_g^{(i)}(f) = \delta_g^{(i+1)}(f). \quad (3.6)$$

The *reconstruction by erosion* can be obtained as its dual operator,

$R_g^\varepsilon(f) = [R_{g^c}^\delta(f^c)]^c$, being f^c the complement image (i.e., the negative). The reconstruction by dilation removes from the reference image g the light objects unconnected with the marker image f and the reconstruction by erosion the dark ones (Figure 3.5).

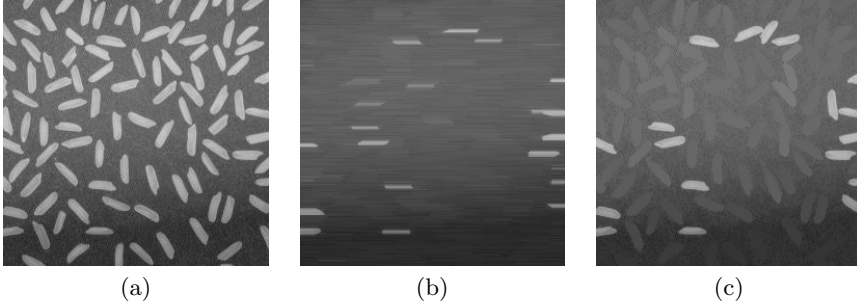


Figure 3.5: Geodesic reconstruction: (a) Reference image, (b) Marker image and (c) Reconstruction by dilation.

Using the *reconstruction by dilation*, the *close-hole operator* can also be defined. For a gray-scale image, it is considered a hole any set of connected points surrounded by connected components of value strictly greater than the hole values. This operator fills all holes in an image f that do not touch the image boundary f_∂ that is used as marker (Figure 3.6):

$$\psi^{ch}(f) = [R_{f^c}^\delta(f_\partial)]^c. \quad (3.7)$$

3.2.2 Stochastic watershed transformation

Watershed transformation is a segmentation technique for gray-scale images (Beucher and Meyer, 1992). This algorithm is a powerful segmentation tool whenever the minima of the image represent the objects of interest and the maxima are the separation boundaries between objects. Due to this fact, the input image of this method

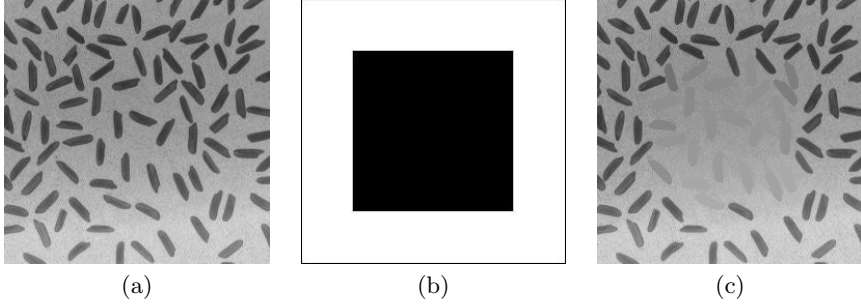


Figure 3.6: *Close-hole operator: (a) Reference image, (b) Marker image and (c) Close-hole.*

is usually a gradient image $\varrho(f)(\mathbf{x})$. The watershed transformation produces a segmentation which can be viewed as: a set of closed contours of segmented regions which will be noted by $WS(\varrho(f))$, or a partition of the space E in a set of classes named $\Pi(WS(\varrho(f)))$.

One problem of this technique is the over-segmentation, which is caused by the existence of numerous local minima in the image normally due to the presence of noise. One solution to this problem is using marker-controlled watershed, $WS(\varrho)_{f_{mrk}}$, in which the markers f_{mrk} artificially indicate the minima of the image. Nevertheless the controversial issue consists in determining f_{mrk} for each region of interest,

$$f_{mrk}(\mathbf{x}) = \begin{cases} 0 & \text{if } \mathbf{x} \in \text{marker} \\ 255 & \text{Otherwise.} \end{cases} \quad (3.8)$$

Note that the number of markers determines the number of regions obtained. The use of few markers along with the existence of borders within the structure to be segmented can also cause that some parts of it are not detected (under-segmentation). So that, the choice of the correct markers is crucial for the effectiveness and robustness of the

algorithm. In marker definition not only markers contained inside the objects of interest (internal markers) are needed but also a marker that determines the image background (external marker). For example, in the case that only a single region was desired, two markers should be used, one internal and other external. Figure 3.50 shows the over and under-segmentation problems related to the watershed transformation when attempting to segment the optic disk.

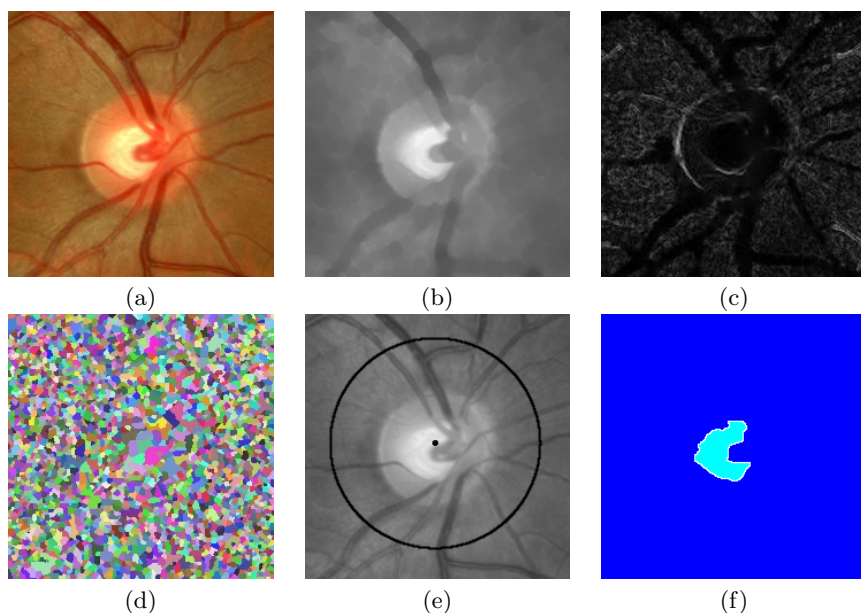


Figure 3.7: *Over and under-segmentation problems of the watershed transformation: (a) Original image, (b) Gray image after closing operator for vessel removal, (c) Gradient image, (d) Classical watershed (over-segmentation), (e) Artificial markers and (f) Marker-controlled watershed using the artificial markers (under-segmentation).*

There exist different strategies to choose the markers to be used in the marker-controlled watershed. The following ones can be stood out:

- Markers based on a seed. This seed can be chosen manually or selected automatically according to the significance of its intensity level, for instance the minima having a local dynamics larger than a given value (Beucher and Meyer, 1992). Another example can be the pixel with the highest/lowest intensity value (Walter et al., 2002).
- Stratified markers. Stratified markers are uniform markers generated within an area. The generation of stratified markers consists of dividing the region to be segmented into a uniform grid, and only the centroid of each region is considered as marker (Chilès and Delfiner, 2012; Bernander et al., 2013).
- Uniform random markers. Random markers are uniformly distributed in the image domain to be segmented (Angulo and Jeulin, 2007).
- Regionalized random markers. Regionalized random markers are non-uniform random markers whose distribution is restricted to areas that accomplished a specific condition, for example they can only be located in high-intensity areas (Angulo and Jeulin, 2007; Angulo and Velasco-Forero, 2010).

In this thesis, stratified and uniform random markers are used. It can be observed in Subsection 3.3.2 and Section 3.5.

Making use of random markers, a watershed transformation variant is used to solve over and under-segmentation problems related to the classical watershed, the stochastic watershed (Angulo and Jeulin, 2007). In this transformation, a given number M of marker-controlled-watershed realizations are performed selecting N random markers in order to estimate a probability density function (*pdf*) of image contours and filter out non-significant fluctuations. Let $\{f_{mrk_i}\}_{i=1}^M$ be M sets of N uniform random markers and $WS_i = WS(\varrho)_{f_{mrk_i}}$ the i th output image of the marker-controlled watershed imposed by f_{mrk_i} . The *pdf*

of image contours is computed by Parzen window method (Duda and Hart, 1973) as follows

$$pdf(\mathbf{x}) = \frac{1}{M} \sum_{i=1}^M (WS_i(\mathbf{x}) * G(\mathbf{x}; s)), \quad (3.9)$$

where $G(\mathbf{x}; s)$ represents a Gaussian kernel of variance σ^2 and mean μ ($\mu = 0$)

$$G(\mathbf{x}; s) = \frac{1}{2\pi\sigma^2} e^{-\left(\frac{\|\mathbf{x}\|^2}{2\sigma^2}\right)}. \quad (3.10)$$

Figure 3.8 illustrates how the *pdf* is generated. Afterwards, it is necessary to perform a last marker-controlled watershed on the *pdf* obtained.

This type of watershed works better than other marker-based watershed transformations used previously in the literature. In the next sections, two different applications of the stochastic watershed will be explained in detail, one for vessel centerline extraction and other for optic disk segmentation.

3.3 Vessel segmentation

Retinal vasculature is able to indicate the status of other vessels of the human body. Indeed, the retina is the only location where blood vessels can be directly visualized non-invasively in vivo (Patton et al., 2006). That is the reason why its study is usually included in the standard screening of any patient with diseases in which the vessels may be altered. In many instances, preclinical signs are not easily recognized and often appear as signs or symptoms that are not specific for a particular disease, but the retina and its blood vessel characteristics

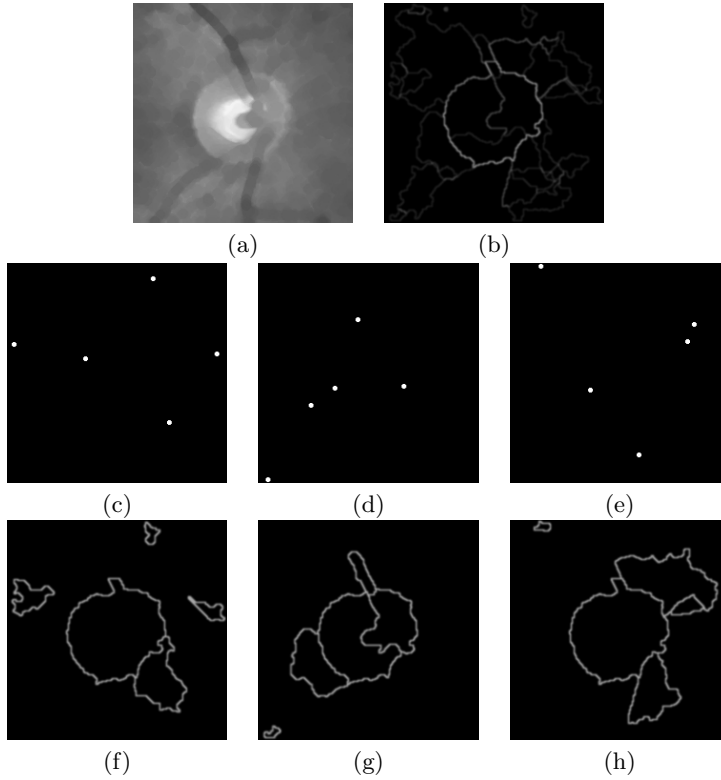


Figure 3.8: *Probability density function: (a) Original image f , (b) Probability density function of contours computed by Parzen window method for $N=5$ and $M=5$, (c-e) Random markers f_{mrk_i} and (f-h) Corresponding marker-controlled-watershed realizations WS_i . The pdf is generated by combining the different WS_i .*

have shown to be a window into several disease processes (Cree and Jelinek, 2011).

In this section, algorithms for vascular network determination will be presented, both for extraction of the complete retinal vasculature and the vessel centerline.

3.3.1 Vessel extraction

The proposed segmentation method is based on mathematical morphology and curvature evaluation for the detection of retinal vascular tree. Although fundus images are RGB images, only monochrome images obtained from the green band are drawn on because they provide an improved visibility of retinal blood vessels as shown in Figure 3.9.

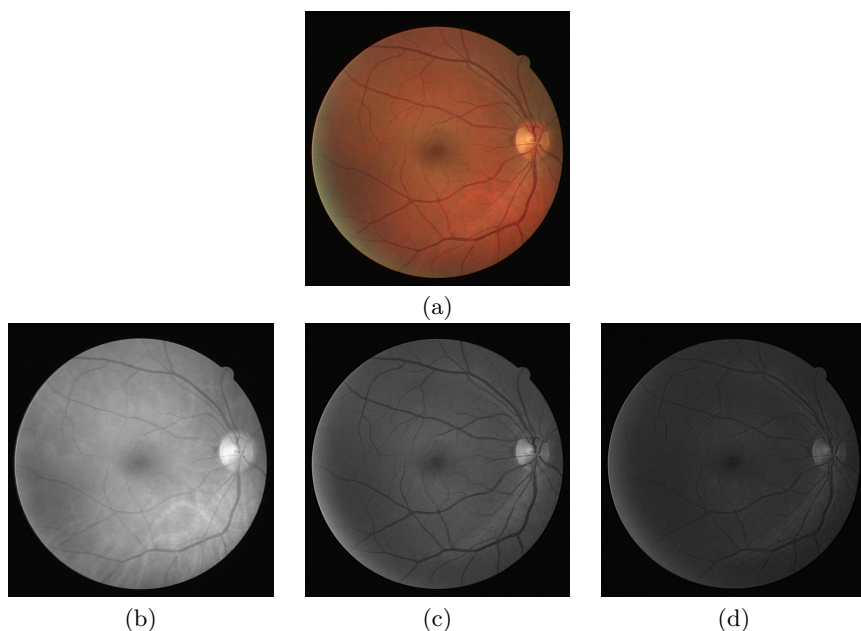


Figure 3.9: *RGB to gray conversion: (a) Original RGB image, (b) Red component, (c) Green component and (d) Blue component.*

Non-uniform illumination and low contrast are typical and inherent problems to the image capture technique. Moreover, in the case of retinal images, both problems are especially pronounced due to the fact that the retina is a spherical structure and it is necessary the use of a spotlight to capture the image. So, they should be corrected previously

to the image processing. In this work, next local transformation for shade correction (Walter and Klein, 2002) was used:

$$\Gamma(f)(t) = \begin{cases} \frac{\frac{1}{2}(u_{max}-u_{min})}{(\mu_f-t_{min})^r}(t-t_{min})^r + u_{min} & \text{if } t \leq \mu_f \\ -\frac{\frac{1}{2}(u_{max}-u_{min})}{(\mu_f-t_{max})^r}(t-t_{max})^r + u_{max} & \text{if } t > \mu_f, \end{cases} \quad (3.11)$$

where t_{min} and t_{max} are the minimum and maximum gray level of the image respectively, u_{min} and u_{max} are the target levels (typically 0 and 255 respectively), μ_f is the mean value of the image for all pixels within a window centered at the current pixel \mathbf{x} of the fixed size L , and the parameter r is used to control the contrast increasing (experimentally $L = 51$ and $r = 3$ for fundus images). The effect of this transformation can be observed in Figure 3.10. Note that all size parameters used in the vessel extraction method are optimized for the resolution of the images belong to DRIVE database (Staal et al., 2004), which will be described in the material subsection (Subsection 3.7.1).

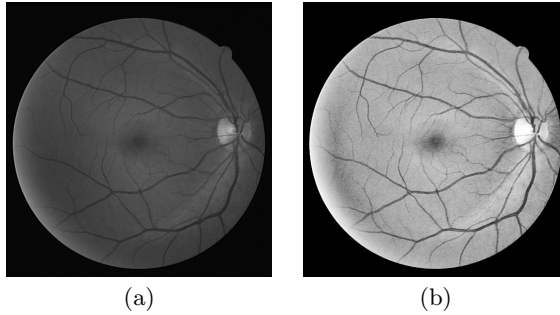


Figure 3.10: Image enhancement: (a) Green component and (b) Local shade correction.

After image pre-processing, the following stages are implemented for vessel segmentation:

First, a small opening, using a disk of radius 1 (B_1) as structuring element (SE), is performed on the enhanced green component image to fill in any gaps of the vessels which could induce errors in segmentation. See Figure 3.11.

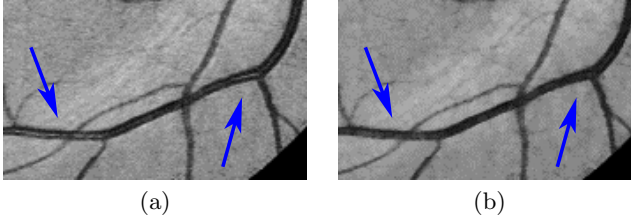


Figure 3.11: *Small opening: (a) Enhanced green component and (b) Opening using B_1 as SE.*

Then, principal curvature is calculated as the maximum eigenvalue (λ_{max}) of the Hessian matrix (Martinez-Perez et al., 2007)

$$H = \begin{pmatrix} \partial_{xx}f & \partial_{xy}f \\ \partial_{yx}f & \partial_{yy}f \end{pmatrix}, \quad (3.12)$$

where $\partial_{ij}f$ represents the second directional derivatives of an image $f(x, y)$. The Hessian matrix is calculated at different scales ($s = \{0, 2, 8, 14\}$) by convolving the original image $f(x, y)$ with a Gaussian kernel G of variance s^2 ,

$$f_s(x, y; s) = f(x, y) \otimes G(x, y; s) = f(x, y) \otimes \frac{1}{2\pi s^2} e^{-\frac{x^2+y^2}{2s^2}}. \quad (3.13)$$

Finally, the principal curvature f_κ is obtained by normalizing each λ_{max}

by $2s$ and computing the local maxima over scales:

$$f_{\kappa} = \max_s \left(\frac{\lambda_{max}(s)}{2s} \right). \quad (3.14)$$

If the principal curvature is directly calculated on the enhanced image, all structures with high curvature are highlighted, not only the vessels. The optic disk border has also high curvature but it should not be detected. This is a typical problem that occurs in most edge detection methods for vessel segmentation. To solve this problem, a dual top-hat, with a circular SE larger than the biggest vessel (B_2), is applied with the goal of extracting all of them and eliminating structures with high curvature that are not vessels, as the optic disk. Figure 3.12 illustrates the difference between calculating the principal curvature directly on the enhanced image and on the result of the dual top-hat.

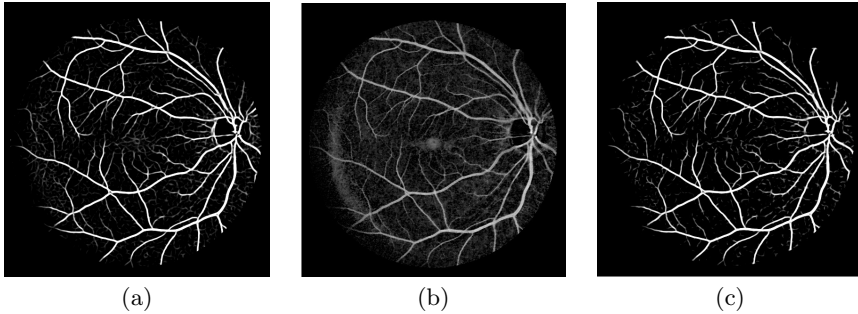


Figure 3.12: *Principal curvature computation: (a) Principal curvature of the enhanced image, (b) Dual top-hat on the enhanced image and (c) Principal curvature of the dual top-hat.*

After that, a reconstruction by dilation is applied to reconstruct the principal curvature from a supremum of openings ($\gamma_B^{sup}(f) = \bigvee_i \{\gamma_{B_i}(f)\}$). In particular, the openings were calculated using a line of size 31 as SE every 15° ($\mathbf{B}_3 = \{B_{3_1}, B_{3_2}, \dots, B_{3_N}\}$, being N the total number of openings). This operation removes any structure smaller

than this SE in any orientation obtaining a cleaner background. See Figure 3.13(a) and 3.13(b).

Finally, to binarize the resulting image a k-means clustering (MacQueen, 1967) is used with a k value equals to 3, giving rise to $\mathbf{f}_{\text{km}} = (f_1, f_2, f_3)$. Afterwards, a modification of the k-means output is carried out; two of the three obtained clusters are defined as vessel ($f_{\text{out}} = f_1 \cup f_2$), considering that f_3 corresponds to the background. Three classes are required because thick and thin vessels can be very different. Figure 3.13(c) depicts the resulting image.

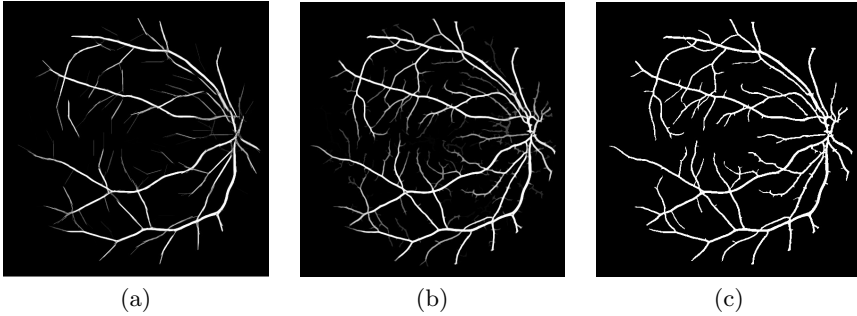


Figure 3.13: *Final vessel segmentation steps: (a) Supremum of openings, (b) reconstruction by dilation and (c) K-means clustering (segmented image).*

Algorithm 3.1 summarizes the main steps of the segmentation method.

The method presented in this Subsection as the same way as the method of Zana and Klein *et al.* (Zana and Klein, 2001) is based on mathematical morphology and curvature evaluation for the detection of the vascular retinal tree. However, the operations used for this purpose are different. Zana and Klein performed a geodesic reconstruction of a supremum of openings to remove noise from the original image while preserving most of the capillaries. Then, a sum of top hats is calculated to improve the contrast of the linear parts. Afterwards, the curvature is computed by means of the Laplacian operator and the final result

Algorithm 3.1: Vessel segmentation

Data: Original RGB fundus image $\mathbf{f} = (f_R, f_G, f_B)$, Scale vector $\mathbf{s} = [0, 2, 8, 14]$, Gaussian kernel G

Result: Vessel binary mask, f_{out}

initialization: B_1, B_2, \mathbf{B}_3 (as main text) ;

$f_{in} \leftarrow f_G$ Green component selection ;

$f_{enh} \leftarrow \Gamma(f_{in})$ Image enhancement ;

$f_{op} \leftarrow \gamma_{B_1}(f_{enh})$ Opening ;

$f_{dth} \leftarrow \rho_{B_2}(f_{op})$ Dual top-hat ;

Principal curvature:

for $i \leftarrow 1$ **to** $length(\mathbf{s})$ **do**

$f_{s_i} \leftarrow f_{dth} \otimes G(s_i)$;

$H_{s_i} \leftarrow H(f_{s_i})$;

$\lambda_{max_{s_i}} \leftarrow \max\{eig(H_{s_i})\}$;

$\lambda'_{max_{s_i}} \leftarrow \frac{\lambda_{max_{s_i}}}{2s_i}$;

end

$f_{\kappa} \leftarrow \max_{\mathbf{s}} (\lambda'_{max_{\mathbf{s}}})$;

$f_{so} \leftarrow \gamma_{\mathbf{B}_3}^{sup}(f_{\kappa})$ Supremum of openings ;

$f_{rec} \leftarrow R_{f_{\kappa}}^{\delta}(f_{so})$ Reconstruction by dilation ;

$f_{out} \leftarrow MKM(f_{rec})$ Modified k-means clustering ;

is obtained with an alternating filter. The proposed method calculates the curvature of a dual top-hat and then a reconstruction by dilation is applied to remove noise. The final result is achieved through a k-means clustering.

3.3.2 Centerline extraction

In general, the detection of retinal vascular network is necessary before analyzing vessel features. The most common approach in the literature is a first stage of vessel segmentation, then the skeletonization

of the detected vessels and finally the analysis of different features on the vascular skeleton as vessel caliber or bifurcation angles. The major drawback of this approach is the dependence of the different stages with the previous ones in addition to computational cost. Based on these facts, this subsection is focused on obtaining the retinal skeleton in a direct way avoiding the segmentation stage. Its goal is to reduce the number of necessary steps in the fundus image processing. As a consequence, this would also reduce the dependency of previous stages. Specifically, the method proposed for this purpose is mainly based on mathematical morphology along with curvature evaluation. Two main steps are involved: in the first step, the principal curvature is calculated on the retinal image. In the second step, the stochastic watershed transformation, presented in Subsection 3.2.2, is applied to extract the vascular skeleton. The main stages are included in the flowchart shown in Figure 3.14.

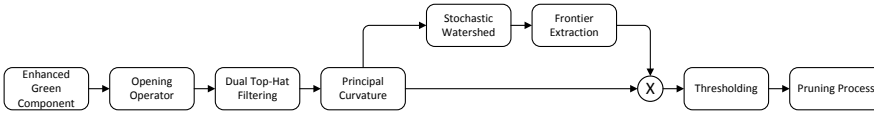


Figure 3.14: *Flowchart for skeleton extraction.*

The first steps of the method are the same that those included in the method for vessel segmentation presented in Subsection 3.3.1. Although fundus images are RGB format (Figure 3.15(a)), only the green component is used (Figure 3.15(b)). Moreover, this gray image is enhanced by means of the Equation 3.11. Then, a small opening, using a disk of radius 1 as SE (B_1), is performed on the enhanced green component image to fill in any gaps in vessels that could provoke subsequent errors, for example due to brighter zone within arteries. Next, a dual top-hat, with a SE larger than the widest vessel (B_2), is applied with the goal of extracting all of them and eliminating structures with high gradient that are not vessels, as occurs with the optic disk border (Figure 3.15(c)). Afterwards, with the aim of highlighting the

vessels on the background, principal curvature, f_κ , is calculated as the maximum eigenvalue of the Hessian matrix (Martinez-Perez et al., 2007) resulting the image shown in Figure 3.15(d).

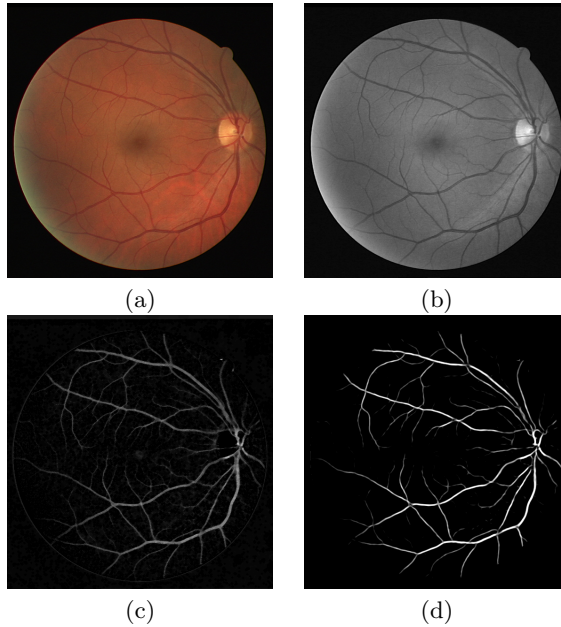


Figure 3.15: First steps of the skeleton extraction process: (a) Original fundus image, (b) Green component, (c) Dual top-hat filtering and (d) Principal curvature.

However, the following steps are different from those of the previous segmentation method and the stochastic watershed is applied to the curvature image. This transformation uses random markers to build a probability density function (*pdf*) of contours (Figure 3.16(a)), which is then segmented by a last marker-controlled watershed. The vascular skeleton is part of the frontiers of the resultant regions as can be observed in Figure 3.16(b).

In that case, both for *pdf* obtaining and for last marker-controlled

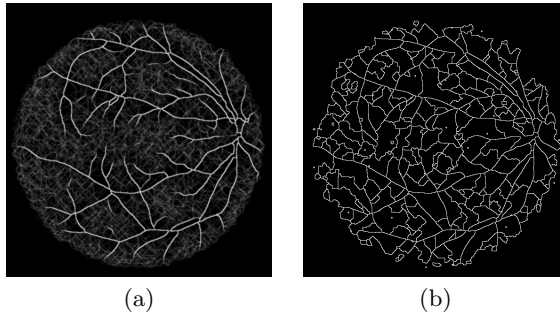


Figure 3.16: Use of the stochastic watershed for vascular skeleton extraction: (a) Probability density function (pdf) of contours obtained with 10 simulations and 300 random markers and (b) Watershed frontiers.

watershed, random markers are combined with some controlled markers. This is due to the morphology of the vascular network which contains multitude of vessel crossings. In addition to the random markers, it is forced that there is one marker at least in the area delimited by the crossing of two vessels (controlled markers). This methodology avoids that the vessels close to some crossing are not detected by the watershed transformation. The crossing areas are determined by means of the residue of the close-hole operator on f_κ and then one or more markers are chosen randomly within these areas giving place to the controlled markers. This problem is illustrated in Figure 3.17, where only a region of interest is shown for better visualization.

In order to discriminate which watershed frontiers are significant and which ones are not and should be filtered out, the frontiers are multiplied by f_κ and then are thresholded (Figure 3.18(a)) using a fixed threshold, experimentally $t = 0.05$. Once the skeleton is obtained, a pruning process is applied to remove possible spurs giving rise to the final result of the method (Figure 3.18(b) and 3.18(c)).

The implemented pruning process is characterized by removing spur branches but without altering the main branches. Only the branches

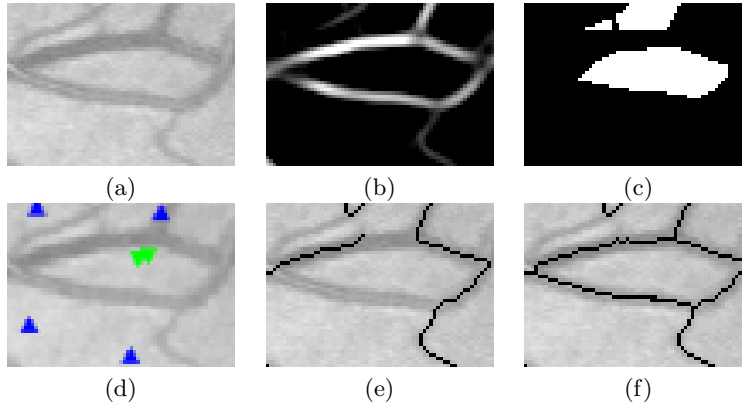


Figure 3.17: *Stochastic watershed on the crossing of two vessels: (a) Enhanced green component, (b) Principal curvature (f_{κ}), (c) Residue of close-hole operator, (d) Random (blue \blacktriangle) and two controlled (green \blacktriangledown) markers, (e) Result of the stochastic watershed using only the random markers shown in blue and (f) Result of the stochastic watershed combining random and controlled markers (blue and green).*

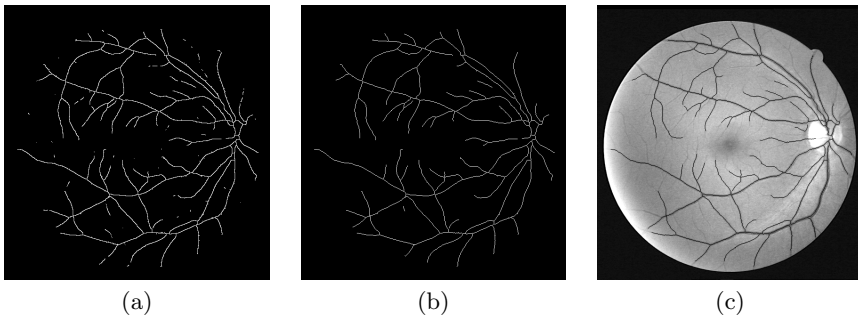


Figure 3.18: *Final steps of the skeleton extraction process: (a) Product between the principal curvature and the watershed frontiers after thresholding, (b) Pruning and (c) Final result.*

whose size is less than a threshold ($n_{max} = 15$) are removed while the other are kept intact (Walter, 2003). The pruning method is based on

defining a function $\Upsilon(S)$ which assigns to each point from the skeleton S the number n of thinning sequences needed to remove it from S . The value of the function $\Upsilon(S)$ is $n_{max} + 1$ for the points $\mathbf{x} \in S$ which are not removed after n_{max} thinning processes. Then, making use of the function $\Upsilon(S)$, it is possible differentiate between the secondary and the main skeleton branches. A branch is considered as secondary if $\Upsilon(S(\mathbf{x}_1)) - \Upsilon(S(\mathbf{x}_2)) > 1$, being \mathbf{x}_1 and \mathbf{x}_2 two adjacent points of the skeleton branch. Afterwards, the secondary branches are disconnected from the main branches and a reconstruction by dilation is applied using this image as reference and being the marker image that defined by Equation 3.15. This operation manages to reconstruct the original skeleton but without spur branches. Figure 3.19 shows an example of the main pruning steps.

$$mrk = \begin{cases} 1 & \text{if } \Upsilon(S(\mathbf{x})) = n_{max} + 1 \\ 0 & \text{otherwise} \end{cases} \quad (3.15)$$

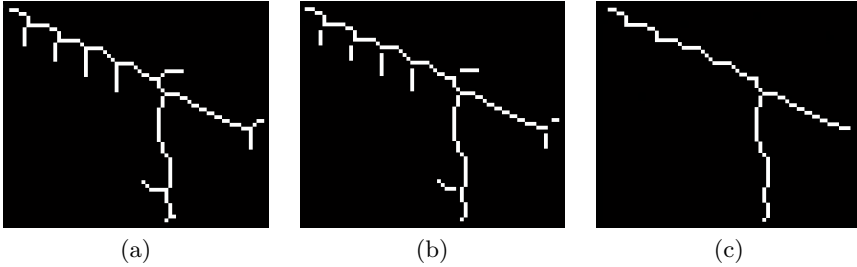


Figure 3.19: *Pruning process: (a) Original skeleton, (b) Skeleton with the secondary branches disconnected and (c) Pruned skeleton.*

Algorithm 3.2 summarizes the steps of the vessel centerline extraction method.

Algorithm 3.2: Vessel centerline extraction

Data: Original RGB fundus image $\mathbf{f} = (f_R, f_G, f_B)$, Scale vector $\mathbf{s} = [0, 2, 8, 14]$, Gaussian kernel G

Result: Vessel centerline, f_{out}

initialization: B_1, B_2 (as main text) ;

$f_{in} \leftarrow f_G$ Green component selection ;

$f_{enh} \leftarrow \Gamma(f_{in})$ Image Enhancement ;

$f_{op} \leftarrow \gamma_{B_1}(f_{enh})$ Opening ;

$f_{dth} \leftarrow \rho_{B_2}(f_{op})$ Dual top-hat ;

Principal curvature:

for $i \leftarrow 1$ **to** $length(\mathbf{s})$ **do**

$f_{s_i} \leftarrow f_{dth} \otimes G(s_i)$;

$H_{s_i} \leftarrow H(f_{s_i})$;

$\lambda_{max_{s_i}} \leftarrow \max\{eig(H_{s_i})\}$;

$\lambda'_{max_{s_i}} \leftarrow \frac{\lambda_{max_{s_i}}}{2s_i}$;

end

$f_\kappa \leftarrow \max_{\mathbf{s}} (\lambda'_{max_{\mathbf{s}}})$;

$f_{ws} \leftarrow WS(f_\kappa)_{f_{mrk}}$ Stochastic Watershed with random and controlled markers ;

$f_{th} \leftarrow (f_\kappa \times f_{ws}) < t$ Thresholding ;

$f_{out} \leftarrow \Upsilon(f_{th})$ Pruning ;

3.4 Significant points

This section presents how to detect one of the main features that define and characterize the retinal vascular network: its significant points. The detection of significant points in the retinal vascular tree will increase the information about the vascular structure allowing its use for medical diagnosis (Calvo et al., 2011).

In general, the significant points of the vascular network are detected on vessel centerline. As mentioned in the previous section, the centerline

can be obtained after a skeletonization process of the vessels previously segmented or through some method by which the skeleton is directly obtained as that presented in Subsection 3.3.2.

In the vascular skeleton there are three types of significant points: terminal, bifurcation and crossing points. All of them must be detected due to their interest to characterize the relations between the different branches of the skeleton, i.e., relations between the parent and daughter branches.

Hit-or-miss transformation (HMT) is a binary morphological operation which is used for detecting specific patterns in an image (Gonzalez et al., 2004). Therefore, it can be applied to detect the significant points on a binary skeleton image. The structuring element (SE) employed in this operation is called composite structuring element since it contains two SE. The first, denoted by B_{FG} , defines the set of pixels that should match the foreground (positive pixel values) while the second, denoted by B_{BG} , defines the set of pixels that should match the background (zero pixel values). By definition, B_{FG} and B_{BG} share the same origin and are disjoint sets, i.e., $B_{FG} \cap B_{BG} = 0$. Depending on whether the origin belongs to B_{FG} or B_{BG} the HMT extracts foreground or background pixels (Soille, 2003). Figure 3.20 depicts a composite structuring element $\mathbf{B} = (B_{FG}, B_{BG})$ where B_{FG} is denoted by 1's, B_{BG} by 0's and the rest of values are ignored. In this example, the HMT would detect the pixels with a neighbor on the left but that up, down and to the right did not have any. The value of the left diagonals does not matter, it could be 0 or 1.

$$\begin{array}{|c|c|c|} \hline & 0 & 0 \\ \hline 1 & 1 & 0 \\ \hline & 0 & 0 \\ \hline \end{array} \quad \mathbf{B} \quad = \quad \begin{array}{|c|c|c|} \hline & & \\ \hline 1 & 1 & \\ \hline & & \\ \hline \end{array} \quad B_{FG} \quad + \quad \begin{array}{|c|c|c|} \hline & 1 & 1 \\ \hline & & 1 \\ \hline & 1 & 1 \\ \hline \end{array} \quad B_{BG}$$

Figure 3.20: Example of a composite structuring element.

The HMT of a set X by a composite structuring element $\mathbf{B} =$

(B_{FG}, B_{BG}) can be written in terms of an intersection of two erosions (Soille, 2003):

$$HTM_B(X) = \varepsilon_{B_{FG}}(X) \cap \varepsilon_{B_{BG}}(X^c), \quad (3.16)$$

where X^c is the complement set (i.e., the negative).

3.4.1 Terminal and bifurcation points

The hit-or-miss transformation can be directly applied to the vascular skeleton to locate terminal and bifurcation points using the different SE shown in Figure 3.21 and 3.22. It must be remembered that B_{FG} is denoted by 1's and B_{BG} by 0's. Note that these SE must be used in all their orientations, one each 90° , so four hit-or-miss iterations are required for each one.

	0	0
1	1	0
	0	0

B_{1_1}

1	0	0
0	1	0
0	0	0

B_{1_2}

Figure 3.21: Structuring elements used for detecting terminal points.

	1	
1	1	
	1	

B_{2_1}

1		
0	1	
1	0	1

B_{2_2}

	1	
	1	
1	0	1

B_{2_3}

	1	
1	1	
		1

B_{2_4}

Figure 3.22: Structuring elements used for detecting bifurcation points.

The choice of the SE shape is related to the topology of the point in question. For example, on the one hand, any terminal point will have only one neighbor and it will be round by background pixels and, on the other hand, a bifurcation point will have three neighbors located in

particular positions. This can be appreciated clearly if the SE defined in Figure 3.22 are observed.

Retinal skeleton is an one-pixel-thick structure fully 8-connected. However, when the significant points belonging to the skeleton are being looked for, it is wanted to avoid the multiple paths that are inherent in this type of connectivity. Therefore, previously to point detection, it is necessary to convert the skeleton from 8-connectivity to m -connectivity so that the multiple paths are removed.

A pixel p has four horizontal and vertical neighbors $N_4(p)$ and four diagonal neighbors $N_D(p)$. All these neighbors are called the 8-neighbors of p , denoted by $N_8(p)$. Two binary pixels p and q are 8-connected if q is in the set $N_8(p)$ but they are m -connected if

1. q is in $N_4(p)$, or
2. q is in $N_D(p)$ and $N_4(p) \cap N_4(q) = \emptyset$.

The difference between 8-connectivity and m -connectivity can be appreciated in Figure 3.23. This conversion is necessary because the central pixel makes that there is not only a possible path in the skeleton therefore the detection of significant points and any tracking process performed later could be erroneous.

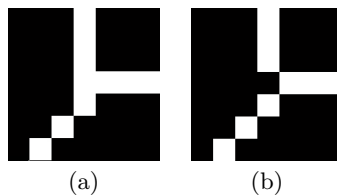


Figure 3.23: *8-connectivity to m -connectivity conversion (a) 8-connected skeleton and (b) m -connected skeleton.*

In the case of the 8-connectivity, the multiple paths are manifested in

four basic patterns which are shown in Figure 3.24. It can be observed that the case shown in Figure 3.23 corresponds to the patterns B_{3_2} and B_{3_3} .

0	1	
1	1	
		0

B_{3_1}

	1	0
	1	1
0		

B_{3_2}

0		
	1	1
	1	0

B_{3_3}

		0
1	1	
0	1	

B_{3_4}

Figure 3.24: Structuring elements used to convert an 8-connected skeleton to m -connectivity.

The HMT allows to detect all these patterns. Then, the central pixels must be changed to 0 for eliminating the multiple paths. The conversion from 8 to m -connectivity can be performed through a basic sequence of morphological steps:

$$\begin{aligned}
 \Theta_1(X, B_{3_1}) &= X - HMT_{B_{3_1}}(X) = X \cap \left[HMT_{B_{3_1}}(X) \right]^c \\
 \Theta_2(\Theta_1, B_{3_2}) &= \Theta_1 - HMT_{B_{3_2}}(Y_1) = Y_1 \cap \left[HMT_{B_{3_2}}(Y_1) \right]^c \\
 \Theta_3(\Theta_2, B_{3_3}) &= \Theta_2 - HMT_{B_{3_3}}(Y_2) = Y_2 \cap \left[HMT_{B_{3_3}}(Y_2) \right]^c \\
 \Theta_4(\Theta_3, B_{3_4}) &= \Theta_3 - HMT_{B_{3_4}}(Y_3) = Y_3 \cap \left[HMT_{B_{3_4}}(Y_3) \right]^c \\
 \Theta(X, \mathbf{B}_3) &= \Theta_4,
 \end{aligned} \tag{3.17}$$

where X is the input image that contains the 8-connected skeleton and Θ the output image with the corresponding skeleton with m -connectivity.

3.4.2 Crossing points

Due to the fact that the intersections between different branches of the vascular tree are formed, usually, by a set of pixels, most crossing points cannot be detected by pattern recognition on skeleton, i.e.,

through the hit-or-miss transformation. It can only be applied in simple crossing point detection using the SE shown in Figure 3.25.

0	1	0		1	0	1
1	1	1		0	1	0
0	1	0		1	0	1
B_{4_1}				B_{4_2}		

Figure 3.25: Structuring elements used for detecting simple crossing points.

However, practically most, if not all crossing points are not simple, or in other words, the branches do not intersect in only one pixel but several points can belong to the same intersection. This provokes that the extremes of the intersection are considered as bifurcation because both of them have three neighbors and accomplishes some of the characteristic pattern of the bifurcation points. Figure 3.26 represents the different types of crossing points. The light gray branch intersects with other two branches drawn in dark gray giving rise to a simple (X) and a complex (O) crossing point marked in black.

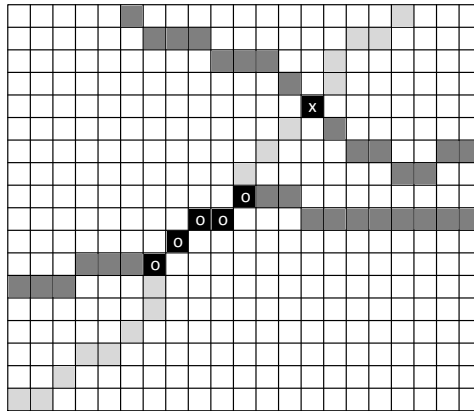


Figure 3.26: Different types of crossing points: simple (X) and complex (O).

Most works of the state of the art consider that the vessel crossing

points are two bifurcation points very close to each other. So, a fixed-size circular window is centered on the candidate bifurcations and if there exist four intersections between the window and the skeleton, the point is marked as crossing. The problem of this approach is that the crossing point detection depends on a large degree on the window size. If the size is too small, the crossings are not detected and if the size is too big other vessels not belonging to this crossing can intersect with the window. Moreover, it must be taken into account that the size of the intersections vary from one case to another.

Retinal vessels have their origin in the optic disk head. From this center, the vessels bifurcate and constitute the retinal vascular tree. It is common that arteries and veins intersect in some occasion and generate the crossing points under consideration (Figure 3.27(a)). This means that when they intersect, as the vessels have a common origin, generate a sort of close loop which will be useful to differentiate if one point is a crossing point or not (Figure 3.27(b)). Based on this idea, a new algorithm is proposed to analyze all points detected initially as bifurcation in order to discriminate those that are really crossing points.

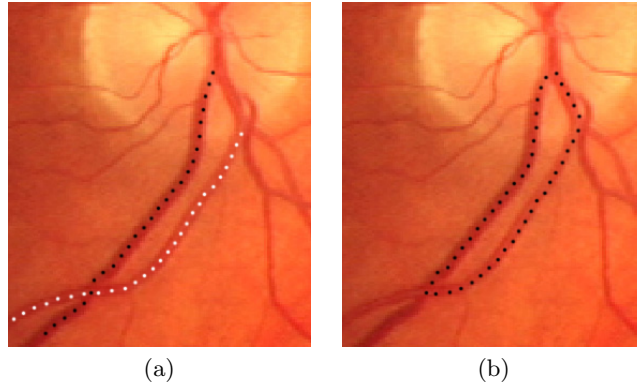


Figure 3.27: *Vessel intersection: (a) Intersection between two vessels and (b) Close loop formed by an intersection.*

First, a bifurcation point is considered as candidate to crossing

point if when a circular window is centered on this point, there are four or more intersections between the skeleton and this window (W). The window radius will be three times the median of the diameter of the vessels in all points detected initially as bifurcation. This size was established empirically. Then, if the candidate point is part of a close loop generated by the skeleton branches, the closest candidate to crossing point is looked for, and if it is directly connected with the previous point and is not part of the same loop, both of them are established as crossing points. Figure 3.28 represents this process. Terminal and bifurcation points detected by the HMT are marked in red and green, respectively (Figure 3.28(a)). In Figure 3.28(b) the candidates to crossing points are highlighted in yellow. The close loops that contain some candidate point are drawn in Figure 3.28(c). Figure 3.28(d) shows the final result with the crossing points detected in white.

With this type of analysis, the more common intersection extremes are completely identified. In addition, it should be taken into account that the pixels between these points are also part of the same intersection.

Algorithm 3.3 summarizes the complete process of the detection of the significant points on the retinal vascular skeleton. $\mathbf{B}_1, \mathbf{B}_2, \mathbf{B}_3, \mathbf{B}_4$ are the composite structuring elements defined in Figures 3.21, 3.22, 3.24 and 3.25.

3.5 Optic disk segmentation

Optic disk (OD) segmentation is a key process in many algorithms designed for the automatic extraction of anatomical ocular structures, the detection of retinal lesions, and the identification of other fundus features. The following examples can be stood out. The OD location helps to avoid false positives in the detection of exudates associated with diabetic retinopathy, since both of them are spots with similar intensity (Walter et al., 2002). The OD margin can be used for establishing standard and concentric areas in which retinal

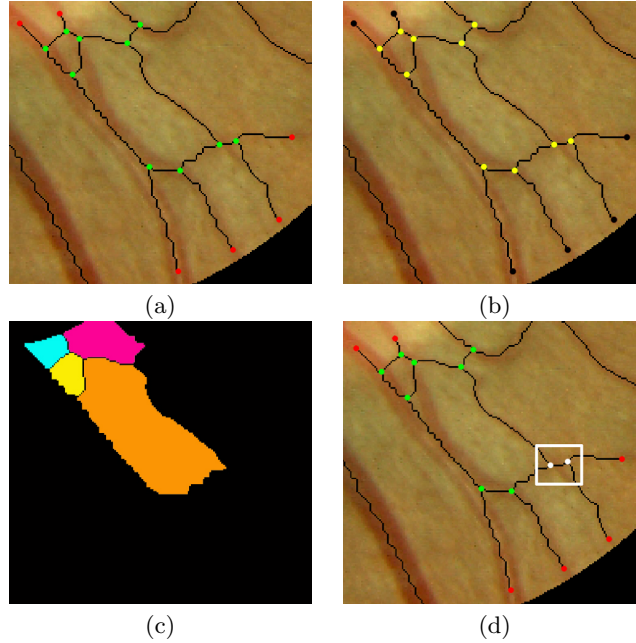


Figure 3.28: Automatic detection of common intersections: (a) Significant points detected by means of HMT (terminal points in red and bifurcation points in green), (b) Crossing point candidates, (c) Close loops formed by branches that contain some candidate point and (d) Crossing points automatically detected (white).

vessel diameter measurements are performed by calculating some important diagnostic indexes for hypertensive retinopathy, such as central retinal artery equivalent (CRAE) and central vein equivalent (CRVE) (Hubbard et al., 1999; Knudtson et al., 2003). The relation between the area of the optic disk and the optic cup is used for glaucoma diagnosis (Liu et al., 2009; Hatanaka et al., 2010). The relatively constant distance between the OD and the fovea is useful for estimating the location of the macula (Niemeijer et al., 2009). The center, or even the border, of the OD also serves as initial point for vessel tracking algorithms due to the fact that all retinal vessels are originated from

Algorithm 3.3: Significant point detection

Data: Image of the retinal skeleton f , Circular window W

Result: Image of the terminal points f_{TP} , Image of the bifurcation points f_{BP} , Image of the crossing points f_{CP}

initialization: $\mathbf{B}_1, \mathbf{B}_2, \mathbf{B}_3, \mathbf{B}_4$ (as main text) ;

$f_m \leftarrow \Theta_{\mathbf{B}_3}(f)$ m -connectivity conversion ;

$f_{TP} \leftarrow HMT_{\mathbf{B}_1}(f_m)$ terminal point detection ;

$f_{BP} \leftarrow HMT_{\mathbf{B}_2}(f_m)$ bifurcation point detection ;

$f_{CP_1} \leftarrow HMT_{\mathbf{B}_4}(f_m)$ simple crossing point detection ;

complex crossing point detection:

for $i \leftarrow 1$ **to** $\sum(f_{BP})$ **do**

if $\sum(W_{BP_i} * f_m) \geq 4$ **then**

if $f_{BP_i} \in \text{close loop}$ **then**

$f_{BP_j} \leftarrow \text{argmin}(f_{BP_i}, f_{BP_j})$;

if (f_{BP_i} is connected with f_{BP_j}) & ($\text{loop}(f_{BP_i}) \neq \text{loop}(f_{BP_j})$) **then**

$f_{CP_{2ij}} = f_{BP_i} + f_{BP_j}$;

end

end

end

end

$f_{CP} = f_{CP_1} + f_{CP_2}$ crossing point detection ;

there (Gagnon et al., 2001).

The automatic method proposed for optic disk segmentation is focused on the stochastic watershed using different operations based on mathematical morphology on a fundus image to obtain the OD-contour. Previously, a pre-processing of the original RGB image is required. The first step of the pre-processing consists of applying PCA to transform the input image to gray scale. This technique combines the most significant information of the three components RGB in a

single image so that it is a more appropriate input to the segmentation method. After segmentation, a post-processing is also performed to fit the final region contour by a circumference. In Figure 3.39, the block diagram of the complete segmentation process is depicted.

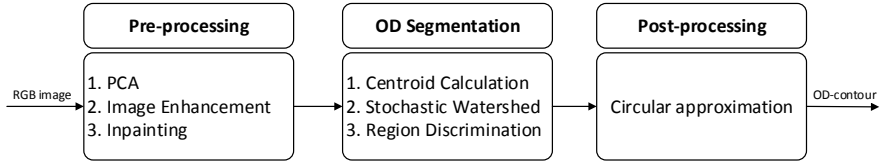


Figure 3.29: *Optic disk segmentation diagram.*

3.5.1 Pre-processing

PCA

Generally, an initial gray-scale image is necessary to carry out most of the segmentation algorithms of the literature. However, in the case of the OD segmentation, each author considers appropriate a different intensity image, such as a band of the original RGB image (Niemeijer et al., 2009; Walter et al., 2002) or a component of other color spaces (Lowell et al., 2004; Osareh et al., 2002). In this work, the use of a new gray-scale image is proposed. Specifically, it is calculated by means of principal component analysis (PCA) because this type of analysis maximizes the separation of the different objects that compose an image so that the structures of the retina are better appreciated. In addition, it is much less sensitive to the existing variability in a fundus image regarding color, intensity, etc.

The central idea of PCA is to reduce the dimensionality of a data set consisting of a number of interrelated variables, while retaining as much as possible of the variation present in the data set. This is achieved by transforming to a new set of variables, the principal components (PCs),

which are uncorrelated, and ordered so that the first few retain most of the variations present in all of the original variables (Jolliffe, 2002).

Suppose that \mathbf{v} is a vector of p random variables with a mean equal to 0, the k th PC, z_k , will be a linear function $\alpha_k' \mathbf{v}$ of the elements of \mathbf{v} where α_k is a vector of p constants $\alpha_{k_1}, \alpha_{k_2}, \dots, \alpha_{k_p}$ and $'$ denotes transpose

$$z_k = \alpha_k' \mathbf{v} = \alpha_{k_1} v_1 + \alpha_{k_2} v_2 + \dots + \alpha_{k_p} v_p = \sum_{j=1}^p \alpha_{k_j} v_j. \quad (3.18)$$

$\alpha_1' \mathbf{v}$ will have maximum variance, $\alpha_2' \mathbf{v}$ will be uncorrelated with $\alpha_1' \mathbf{v}$ and will have maximum variance, and so on, so that at the k th stage $\alpha_k' \mathbf{v}$ will have maximum variance subject to being uncorrelated with $\alpha_1' \mathbf{v}, \alpha_2' \mathbf{v}, \dots, \alpha_{k-1}' \mathbf{v}$.

For a three-channel image transforming to a principal component space creates three new channels in which the first (the most significant) contains the most structural contrast and information. The rank for each axis in the principal set represents the significance of that axis as defined by the variance in the data along that axis. Thus, the first principal axis is the one with the greatest amount of scatter in the data and consequently the greatest amount of contrast and information, while the last principal axis represents the least amount of information such as noise and image artefacts (Russ, 2007). In this case, the PCs are given by

$$z_k = \alpha_k' \mathbf{f} = \alpha_{k_R} (f_R - \bar{f}_R) + \alpha_{k_G} (f_G - \bar{f}_G) + \alpha_{k_B} (f_B - \bar{f}_B), \quad (3.19)$$

where $\mathbf{f}(\mathbf{x}) = (f_R(\mathbf{x}), f_G(\mathbf{x}), f_B(\mathbf{x}))$ represents an RGB image, \bar{f}_i is the mean of each RGB component and $k \in \{1, 2, 3\}$.

Specifically, principal-component axes (α_k) will be the eigenvectors of the covariance matrix Σ . This is the matrix whose (i, j) th element is the covariance between the i th and j th elements of \mathbf{f} when $i \neq j$, and the variance of the j th element of \mathbf{f} when $i = j$ according to

$$\Sigma = \begin{pmatrix} \sigma_{RR} & \sigma_{RG} & \sigma_{RB} \\ \sigma_{GB} & \sigma_{GG} & \sigma_{GB} \\ \sigma_{RB} & \sigma_{GB} & \sigma_{BB} \end{pmatrix}. \quad (3.20)$$

Being:

$$\sigma_{mn} = \frac{1}{N} \sum_{i=1}^N (f_{mi} - \bar{f}_m)(f_{ni} - \bar{f}_n), \quad (3.21)$$

where N is the number of pixels of the image, (m, n) represent the possible combinations between the RGB components and \bar{f}_m and \bar{f}_n determine the mean of the corresponding component, respectively.

Therefore, z_1 , z_2 , and z_3 are the PCs of the image \mathbf{f} so that α_1 is the eigenvector corresponding to the largest eigenvalue of Σ , α_2 corresponding to the second largest eigenvalue, and α_3 corresponding to the smallest one. Figure 3.30 shows the PCs of a RGB fundus image where can be appreciated that the first PC contains the most structural contrast and information.

The first PC is the most significant so, z_1 is chosen as the input image of the method. It must be stressed that to ensure that z_1 contains the most structural contrast and information of the original RGB channels, it should be verified that the largest eigenvalue represents at least a 90% of the total sum of eigenvalues. In spite of this situation is not often in the consulted databases, if the largest eigenvalue represents less than 90%, the components whose sum of eigenvalues get the 90% must be processed in parallel. Afterwards, the input of the watershed transformation will be the pixel-wise maximum of the gradient images.

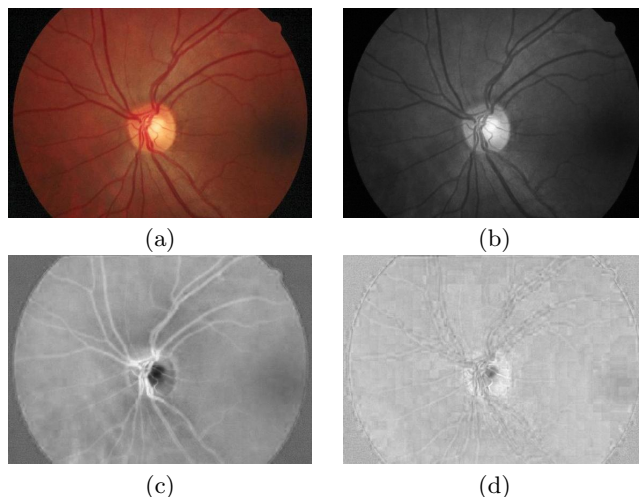


Figure 3.30: *Principal components (PCs): (a) Original fundus image, (b) First PC z_1 , (c) Second PC z_2 and (d) Third PC z_3 .*

The non-uniform illumination of this gray image is corrected by means of the local transformation (Γ) of the Equation 3.11. Γ is applied on z_1 resulting in an enhanced image

$$z'_1 = \Gamma(z_1). \quad (3.22)$$

Inpainting

Retinal vessels are originated from the OD therefore numerous vessels cross its border which makes its discrimination difficult. So, vessel removal of the enhanced image is implemented by an inpainting technique.

Inpainting algorithms are used in diverse applications, from the restoration of damaged photographs to the removal/replacement of selected objects (Guillemot and Le Meur, 2014). These algorithms

usually try to fill selected parts of an image by propagating external information so that structure continuity is preserved.

Although there exist numerous inpainting techniques (Liang et al., 2001; Levin et al., 2003; Telea, 2004; Mairal et al., 2008), a simple diffusion-based algorithm was implemented in this work. Let a binary image $\Omega(\mathbf{x})$ stand for the region to be inpainted and $\partial\Omega$ for its boundary. For each $\partial\Omega$ -pixel, \mathbf{x} , the inpainted-pixel value is computed as

$$P(\mathbf{x}) = \frac{\sum_{k=1}^n \frac{P_k(\mathbf{x})}{l_k}}{\sum_{k=1}^n \frac{1}{l_k}}, \quad (3.23)$$

where P_k denotes the pixel values in a 5 x 5 neighborhood of the pixel under consideration, n is the number of neighboring pixels, and l_k is the distance between the pixel \mathbf{x} and each neighboring pixel. So that the inpainted image $\Psi(f, \Omega)(\mathbf{x})$ of a gray image $f(\mathbf{x})$ is

$$\Psi(f, \Omega)(\mathbf{x}) = \begin{cases} f(\mathbf{x}) & \text{if } \partial\Omega(\mathbf{x}) = 0 \\ P(\mathbf{x}) & \text{if } \partial\Omega(\mathbf{x}) = 255. \end{cases} \quad (3.24)$$

After filling $\partial\Omega$ with the computed values, the $\partial\Omega$ -pixels are removing from Ω and $\partial\Omega$ is recalculated. The process is repeated until the mask is empty and all pixels are filled.

In particular, the region to be inpainted Ω is a binary mask of the retinal vessels, which must have been segmented previously. The method used for vessel segmentation is that proposed in Subsection 3.3.1. Afterwards, a unitary morphological dilation of the segmented vessels yields the final vessel mask. The purpose of this operation is to make sure that the vessels will be contained in the mask.

The aim of applying the inpainting is to extract the OD-boundary more precisely and to reduce the existing borders within the OD which increase the risk of under-segmentation. This operation gives place to

$$z_1'' = \Psi(z_1', \Omega), \quad (3.25)$$

being z_1' the enhanced image (Equation 3.22), Ω the binary image with the segmented vessels and z_1'' the inpainted image.

In Figure 3.31 the pre-processing of the original image, shown in Figure 3.30(a), can be observed, both the obtaining of the first PC and its enhancement by means of non-linear transformation Γ and also the vessel removal. Note that the whole image is processed although only a region of interest is shown for better visualization.

3.5.2 Processing

Marker selection

As mentioned above, the segmentation method makes use of the stochastic watershed. This transformation uses random markers to build a probability density function (*pdf*) of contours, according to Equation 3.9, which is then segmented by volumic watershed to define the most significant regions. However, in the marker definition not only internal markers (that specify what is the object of interest) are needed, but also an external marker which limits the area to be segmented.

On the one hand, the chosen external marker, f_{ext} , will be a circle of constant diameter centered on the centroid of the optic disk. The size of the circle is related to the image size, so that it is approximated by a 15% of the size of the fundus image (with this size we made sure that the optic disk is included). The challenge is to be able to find the OD centroid prior to being segmented. For this purpose, we work with the centroid of a gray-level image.

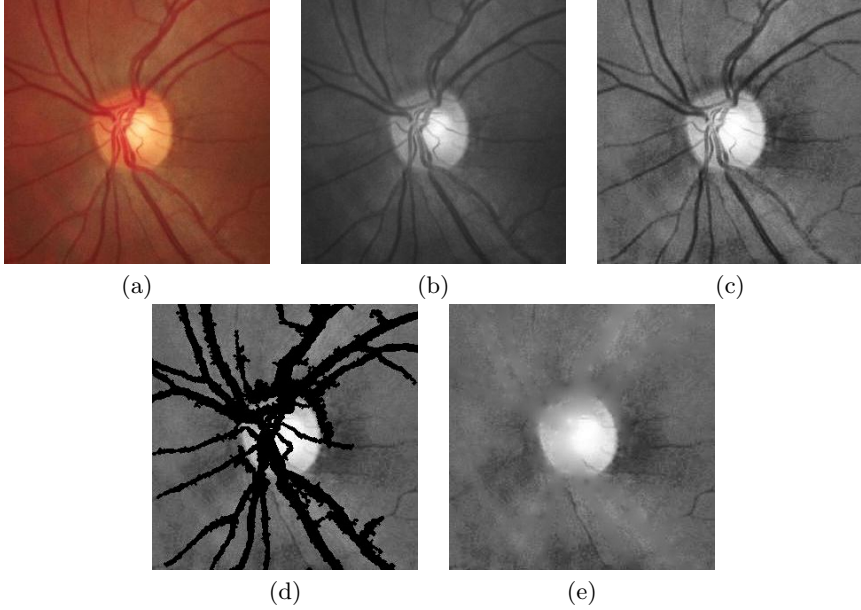


Figure 3.31: *OD pre-processing: (a) Original image, (b) First PC z_1 , (c) Enhanced image z'_1 , (d) Vessel mask to be inpainted Ω , and (e) Inpainted image z''_1 .*

The centroid of a gray-level image can be calculated based on the generalized distance function (GDF) (Vincent, 1998). This algorithm is focused on modifying the classic two-pass sequential distance function (Rosenfeld and Pfaltz, 1968) so that: (1) edge cost is taken into account; (2) raster and anti-raster scans of the image are iterated until stability. Let $N^+(p)$ (resp., $N^-(p)$) be the 8-connected neighborhood of pixel p scanned before p (resp., after p) in a raster scan, and $C_f(p, q) = f(p) + f(q)$ the associated cost to two neighboring pixels p and q , the algorithm of GDF to set X in an image f proceeds as follows:

- Initialize result image d_f : $d_f(p) = 0$ if $p \in X$ and $d_f(p) = +\infty$

otherwise.

- Iterate until stability, for each pixel p :
 - Scan image in raster order:

$$d_f(p) \leftarrow \min\{d_f(p), \min\{d_f(q) + C_f(p, q), q \in N^+(p)\}\}.$$
 - Scan image in anti-raster order:

$$d_f(p) \leftarrow \min\{d_f(p), \min\{d_f(q) + C_f(p, q), q \in N^-(p)\}\}.$$

More specifically, the gray-level centroid of an image f can be obtained as the maximum of the GDF to the image border set B as follows:

- Compute GDF to set B in image f , $d_f(B)$.
- Find the maximal value of $d_f(B)$, u_{max} .
- Threshold $d_f(B)$ with the u_{max} calculated to define set C as image centroid.
- If C has more than one pixel, compute the centroid of set C as the maximum edge distance.

This way for calculating the gray-image centroid combines the centrality of the image with respect to edge distance (i.e. purely geometric) but penalizing this distance in relation to the intensities. Thus, note that it cannot be defined as the center of mass of the intensities or as the center of the brighter and larger zone, since the two effects are combined. In Figure 3.32 the result of applying the algorithm on the first principal component of the original image after inpainting technique can be appreciated.

If Figure 3.32(c) is observed, we can see that the image centroid does not match with the OD centroid. To obtain the desired result, the optic disk should be a brighter zone than the background (sufficiently

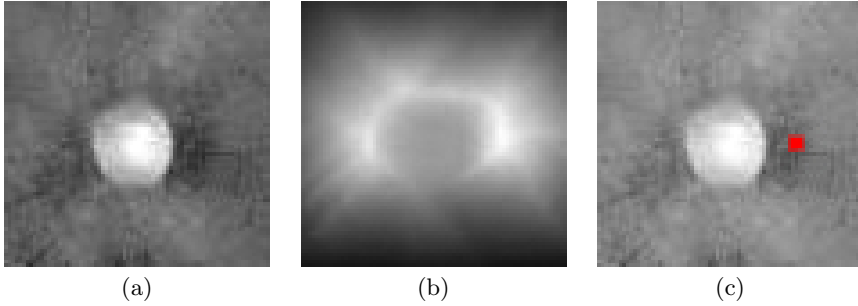


Figure 3.32: *Gray-image centroid on z_1 : (a) First PC (z_1) after inpainting technique, (b) GDF of (a), and (c) Image centroid.*

contrasted) and should have a significant size with respect to the image size. So, to accomplish this condition, the algorithm should be computed on the residue of a close holes of the original gray image as shown Figure 3.33.

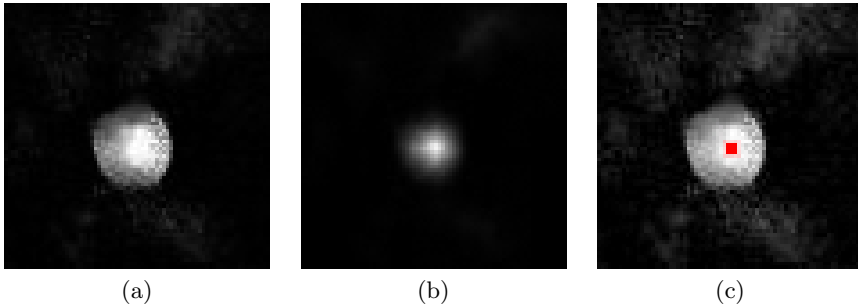


Figure 3.33: *Gray-image centroid on the residue of close operator: (a) Residue of close operator of Figure 3.32(a), (b) GDF of (b), and (c) Image centroid.*

On the other hand, the internal markers, f_{int} , will be uniform random markers generated within the area limited by f_{ext} . See Figure 3.34. Hence, the final set of markers f_m (external and internal) is the

logical OR of both of them,

$$f_m = f_{int} \vee f_{ext}. \quad (3.26)$$

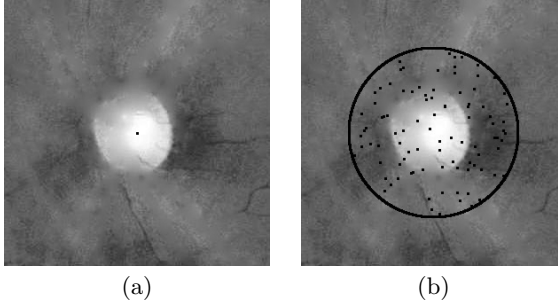


Figure 3.34: Final markers: (a) Centroid image and (b) Uniform random markers limited by an external circle centered on the image centroid.

Stochastic watershed transformation

In particular, the *pdf* is built from 15 marker-controlled-watershed realizations, as showed in Subsection 3.2.2, using as input the gradient of the inpainted image $\varrho(z_1'')$ (Figure 3.35(a)). Therefore, in that case, the *i*th watershed output $WS_i = WS(\varrho(z_1''))_{f_{m_i}}$, being f_{m_i} the final set of markers used in the corresponding simulation. f_{int} is generated for each simulation while f_{ext} is the same in all of them. The number of internal markers used is $N = 100$.

Obtaining a *pdf* of the contours of the watershed regions (Figure 3.35(b)) facilitates the final segmentation, providing robustness and reliability since the arbitrariness in choosing the markers is avoided. Afterwards, the *pdf* can be combined with the initial gradient in order to reinforce the gradient contours which have a high probability resulting

a probabilistic gradient (Angulo and Jeulin, 2007),

$$\rho(z_1'')(\mathbf{x}) = (1 - \lambda)\varrho(z_1'')(\mathbf{x}) + \lambda pdf(\mathbf{x}), \quad (3.27)$$

with $\lambda = 0.5$ the results are in general satisfactory.

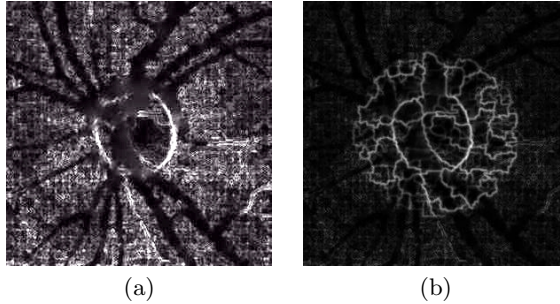


Figure 3.35: Probability density function: (a) Gradient image $\varrho(z_1'')$ and (b) Probability density function of contours using 15 simulations and 100 internal markers.

Finally, a last marker-controlled watershed is applied to $\rho(z_1'')(\mathbf{x})$ using a new set of markers $f_{m'}$, giving rise to the image $\Pi(W S(\rho(z_1'')_{f_{m'}}))$, which is shown in Figure 3.36(b), where the different obtained regions Π can be appreciated. In this case, stratified markers are employed instead of random markers as shown Figure 3.36(a). The reason for using this type of markers is to make sure that every pixel within the area in question belongs to a watershed region. Due to the fact that there are markers located outside the OD, not all regions obtained by the watershed transformation are wanted. If each region of $\Pi(W S(\rho(z_1'')_{f_{m'}}))$ is named as Π_i , the next stage of the proposed method is to discriminate which Π_i are significant and which ones are not and should be filtered out.

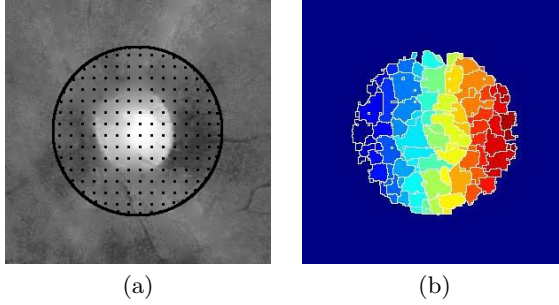


Figure 3.36: Last marker-controlled watershed: (a) Stratified markers, and (b) Watershed regions $\Pi(W S(\rho(z_1'')_{f_{m'}}))$.

Region discrimination

The discrimination between the significant and non-significant regions is based on the average intensity of the region. The value of each region will be equal to

$$\mu_{\Pi_i} = \frac{1}{N_i} \sum_{\mathbf{x} \in \Pi_i} z_1''(\mathbf{x}), \quad (3.28)$$

being N_i the number of pixels of the corresponding region Π_i and z_1'' the inpainted image. Figure 3.37(a) represents the image μ_{Π} where the intensity of each region is equal to μ_{Π_i} . The regions belonging to the optic disk will be light regions around darker regions therefore the residue of a close-hole operator is calculated on μ_{Π} to obtain the regions that accomplish this condition (Figure 3.37(c)). Afterwards, a threshold is applied on the resulting image to select the valid regions. This operation leads to the final OD segmentation (Figure 3.37(d)). The value of the threshold is $u = m - 2s$, being m and s the mean and the standard deviation of the residue of the close-hole operator.

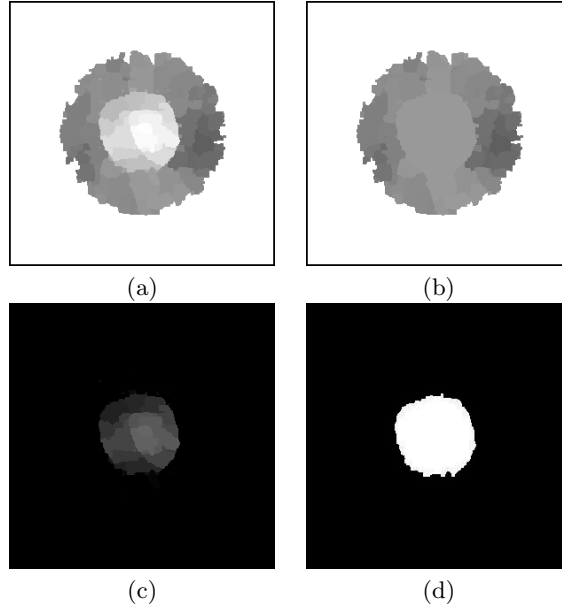


Figure 3.37: *Region discrimination: (a) Average intensity of the watershed regions μ_{Π} , (b) Close-hole operator, (c) Residue of close-hole operator, and (d) Thresholding.*

3.5.3 Post-processing

Once the region of interest is obtained, the result must be fitted to eliminate false contours, which are detected due generally to the blood vessels that pass through the OD. The inpainting technique was performed to remove most of them, as previously mentioned, however some irregularities can still be appreciated in the final region contour (Figure 3.38(a)).

In this work, the OD-contour is estimated as a circle in the same way that in (Aquino et al., 2010; Niemeijer et al., 2009; Lalonde et al., 2001) although an elliptical shape could also be chosen. The main reason to fit the OD by a circle is because this algorithm will

later be used to establish a zone of the retina concentric to the OD-margin according to a standard protocol with the aim to perform vessel diameter measurements (Wong et al., 2004). The fit is performed by means of Kasa's method (Corral and Lindquist, 1998) which lets calculate the center and the radius of the circle that better is adapted to a binary region through least squares. Figure 3.38 shows the contour of the final region obtained by the proposed method and its circular approximation.

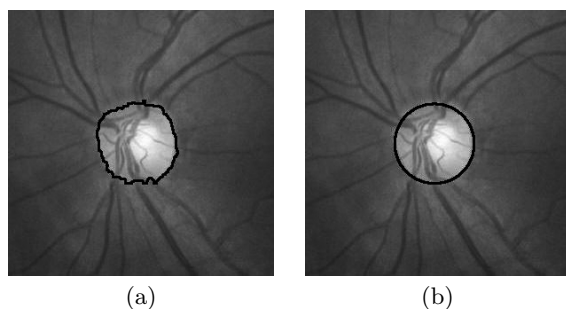


Figure 3.38: *OD post-processing: (a) Contour of the obtained region, and (b) Circular approximation of the OD contour.*

3.6 Application to pathology detection: Hypertensive risk

Most attempts to automate the process of interpretation of retinal vascular imaging are focused on a specific disease, diabetic retinopathy, a disease of high incidence and a significant risk of blindness that occupies a very important part of the medical-surgical activity of the ophthalmologic resources (Goatman et al., 2011; Abramoff et al., 2010). In some of these studies, it was possible to relate the evolution of the disease and the positive or negative response to treatment with retinal vessel caliber (Nguyen et al., 2007; Rogers et al., 2008; Cheung and Wong, 2008). However, there is not too much experience in the use of

these methods to evaluate other types of vascular pathology although fundus photography allows also the determination of cardiovascular risk factors (Abràmoff et al., 2010). Vascular changes produced in systemic diseases usually induce particular modifications in the vessels, such as changes in the angle of intersection between arteries and veins, and changes in the vessel calibers, but the existing solutions only focus on determining the caliber changes (Ruggeri et al., 2007; Muramatsu et al., 2010; Tramontan et al., 2008; Niemeijer et al., 2011; Ortega et al., 2010). Based on these facts, a system capable of detecting the retinal vessels and measuring several geometrical properties from a fundus image was developed. The goal of this software is to assist ophthalmologists in diagnosis and disease prevention, helping them to establish objective relations between the different vessels, to determine cardiovascular risk or other diseases where the vessels can be altered, as well as to monitor the pathology progression and response to different treatments. This tool was applied in a clinical study in order to evaluate sensitivity, specificity and reproducibility of the developed system to discriminate between a normal vascularization and cardiovascular pathology in contrast to the opinion of an expert ophthalmologist obtained through visual inspection of the fundus image.

The computer-aided diagnosis software presented in this section is based on digital fundus image processing for the determination of hypertensive risk. In general, systemic diseases produce vascular changes in our blood system and usually induce specific modifications in the retinal vessels; hence, the system is focused on vessel detection and the measurement of some of their features to characterize any retinal morphological change. So, the main stages involved in the system are: firstly, vessel segmentation process; secondly, retinal tree labelling to identify the pixels that correspond to each vessel and the existing relationship between the branches; and thirdly, performing characteristic measurements on the retinal tree, using the previous information, to quantify significant changes in the vascular network. These data will be used to classify an image as belonging to a healthy subject or to other with cardiovascular risk. The mentioned stages can

be observed in the block diagram of the Figure 3.39 where the entire image processing carried out by the proposed system is illustrated.

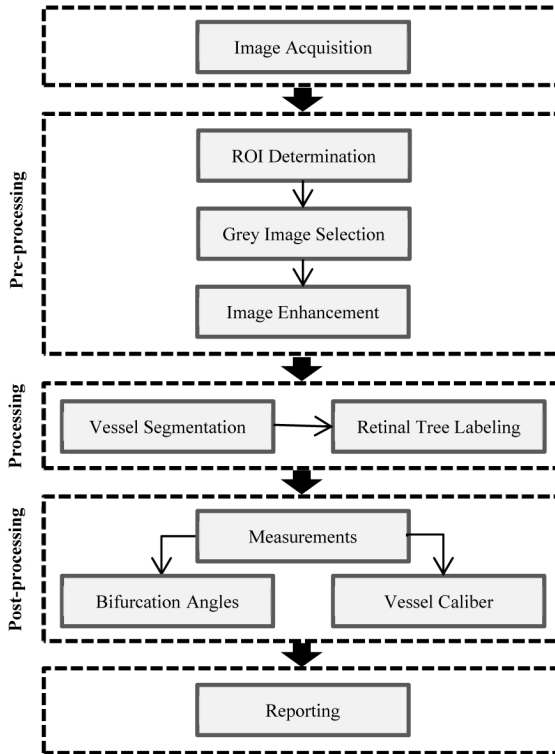


Figure 3.39: Block diagram of the presented tool.

3.6.1 Pre-processing

Generally, based on a standard protocol, the measurement of retinal vessel calibers is focused on a specific region of interest (ROI) of the fundus image (Wong et al., 2004; Knudtson et al., 2003; Hubbard et al., 1999). This area is concentric to the optic disk and it is related to its diameter. So, for that reason, in first place, the proposed system detects

the optic disk (OD) in an automatic way in order to be able to determine the ROI where all measures will be performed. The method used for the extraction of the optic disk is explained in the previous section (Morales et al., 2013).

Once the ROI is established, although original fundus images are RGB images, the system is only drawn on the green band because this band provides an improved visibility of retinal blood vessels. Afterwards, an image enhancement (Walter and Klein, 2002) is applied to improve even more, if it is possible, their visibility (Equation 3.11).

3.6.2 Processing

The segmentation method used by the system is based on mathematical morphology, curvature evaluation and k-means clustering for the detection of a vascular tree (Morales et al., 2012) that is explained in Subsection 3.3.1.

After vessel detection, the vessels must be labelled. Retinal vascular tree labelling is focused on obtaining the skeleton of the vascular tree, detecting significant points and a tracking process (Morales et al., 2012). They are necessary steps to perform later the desired measures.

The skeleton of the vascular tree is obtained by a thinning process from the segmented binary image (Soille, 2003). Next, a pruning process is applied to eliminate possible spurs (Walter, 2003). Significant points (terminal, bifurcation and crossing points) are detected by the method proposed in Section 3.4.

The tracking purpose is to analyze the direction of every pixel of the skeleton and to identify which pixels belong to each branch. The method used is proposed in (Martinez-Perez et al., 2002). The algorithm is able to specify the direction of any skeleton point from a starting point and, therefore, it lets identify the parent and daughter branches.

Note that the method used for vessel segmentation and the later

skeletonization could be replaced by the algorithm for vessel centerline extraction presented in Subsection 3.3.2.

3.6.3 Post-processing

Certain geometric measurements of blood vessels can help to establish whether they have undergone morphological changes over time and facilitate disease diagnosis. The remaining parameters were chosen due to the fact that they have particular interest for the early hypertension detection:

Bifurcation angle: Angle formed by the daughter branches for each bifurcation point. The branches are fitted for straight lines by least-squares into a circular window centered on these points and then the angle α between them is calculated as

$$\alpha = \arccos \left(\frac{|u_1 \cdot v_1 + u_2 \cdot v_2|}{\sqrt{u_1^2 + u_2^2} \cdot \sqrt{v_1^2 + v_2^2}} \right), \quad (3.29)$$

being $\mathbf{u} = (u_1, u_2)$ and $\mathbf{v} = (v_1, v_2)$ the direction vectors of these lines. See Figure 3.40 and 3.41.

Vessel caliber: It is estimated as two times the average of the distance transform value (Soille, 2003) on the skeleton points of the corresponding vessel. The distance transform is an operator applied to binary image. The result of the transform is a gray-level image whose intensity represents the distance from each pixel to the closest boundary. So that the caliber of a vessel V_i is

$$C_{V_i} = \frac{1}{N} \sum_{p \in S_i} d_{V_i}(p), \quad (3.30)$$

being d_{V_i} the distance transform of the vessel V_i , S_i the skeleton of this

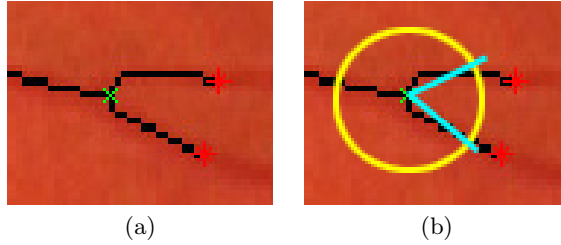


Figure 3.40: *Bifurcation angle measurement: (a) Vessel bifurcation, and (b) Measurement process. A circular window (yellow) is centered on the bifurcation point (green cross) and the daughter branches are fitted for straight lines (cyan). The angle between these lines is directly calculated from their direction vectors.*

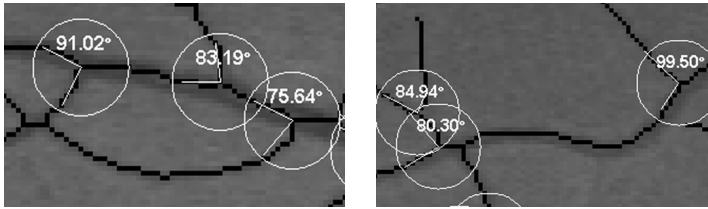


Figure 3.41: *Bifurcation angles measured on two different image areas.*

vessel and N the number of pixels of S_i . Figure 3.42 shows a vessel binary mask and its corresponding distance transform.

As a summary, Figure 3.43 includes the main stages of the image processing performed by the developed software: ROI determination, vessel segmentation and angle and caliber measurement.

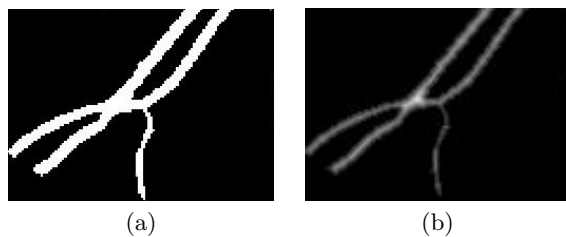


Figure 3.42: *Vessel caliber measurement: (a) Vessel binary mask, and (b) Distance transform.*

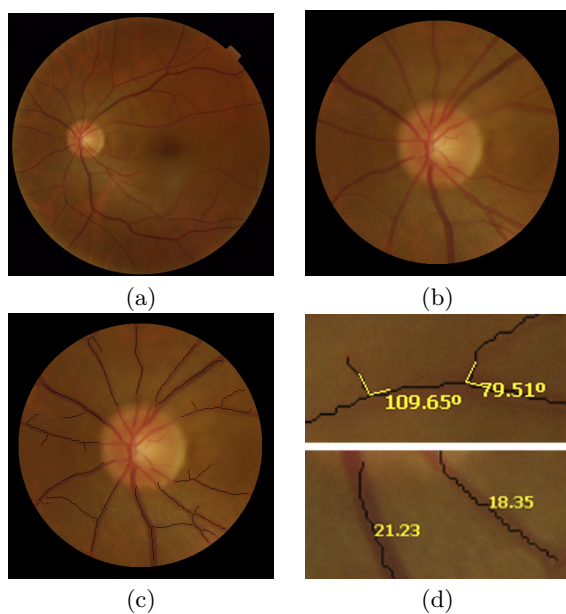


Figure 3.43: *Image processing performed by the developed software: (a) Original fundus image, (b) ROI determination, (c) Vessel segmentation, (d) Bifurcation angles (up) and vessel calibers (down).*

3.7 Results

In this section, the performance of the methods presented in this chapter is detailed along with a description of the databases used for that purpose.

3.7.1 Material

Different public and private databases were used to validate the methods proposed in this chapter. Next, a description of each of them is included.

DRIVE

DRIVE (Digital Retinal Images for Vessel Extraction) ([Staal et al., 2004](#)) is a public database established to enable comparative studies on retinal vessel segmentation.¹ The database is formed by 40 retinal images (768 x 584) belonging to diabetic subjects between 25-90 years of age. 33 do not show any sign of diabetic retinopathy and 7 show signs of mild early diabetic retinopathy.

For each image, a mask image that delineates the field of view is provided as well as manual segmentations of the blood vessels. The images are divided into a training and a test set, both of 20 images. For the training images, only a single manual segmentation is available. For the test cases, two manual segmentations are included.

STARE

STARE (STructured Analysis of the Retina) database ([Hoover et al., 2000](#)) is designed for blood vessel segmentation.² This dataset consists

¹www.isi.uu.nl/Research/Databases/DRIVE/

²www.ces.clemson.edu/~ahoover/stare/probing/index.html

of 20 retinal images (700 x 605). For each image, vessel network is hand labelled by two experts.

DRIONS

DRIONS (Digital Retinal Images for Optic Nerve Segmentation) Database (Carmona et al., 2008) is a public database for benchmarking optic nerve head segmentation from retinal images.³ The database consists of 110 retinal images (600 x 400) whose optic disk is manually segmented by two specialists. The mean age of the patients was 53.0 years (S.D. 13.05), with 46.2% male and 53.8% female and all of them were Caucasian ethnicity. 23.1% patients had chronic simple glaucoma and 76.9% eye hypertension. Some of the 110 images contain visual characteristics related to potential problems that may distort the detection process of the contour of the optic disk: light artefacts (3 images), some of rim blurred or missing (5 images), moderate peripapillary atrophy (16 images), concentric peripapillary atrophy/artefacts (20 images), and strong pallor distractor (6 images).

DIARETDB1

DIARETDB1 database (Kauppi et al., 2007) is a public database for diabetic retinopathy detection.⁴ The database is formed by 89 color fundus images (1500 x 1152). 84 of them contain at least mild nonproliferative signs (microaneurysms) of the diabetic retinopathy, and five are considered as normal which do not contain any signs of the diabetic retinopathy according to all experts who participated in the evaluation. Four medical experts marked the areas related to the microaneurysms, hemorrhages, and hard and soft exudates.

³www.ia.uned.es/~ejcarmona/DRIONS-DB.html

⁴www.it.lut.fi/project/imageret/diaretdb1/

MESSIDOR

MESSIDOR database, kindly provided by the Messidor program partners, was established to facilitate studies of diabetic retinopathy detection.⁵ It contains 1200 eye fundus color images of the posterior pole. The images were captured using 8 bits per color plane at different resolutions (1440 x 960, 2240 x 1488 and 2304 x 1536). 800 images were acquired with pupil dilation and 400 without dilation. For each image, two diagnosis were provided: retinopathy grade and risk of macular edema.

ONHSD

ONHSD (Optic Nerve Head Segmentation Dataset) contains 99 fundal images (640 x 480) taken from 50 patients randomly sampled from a diabetic retinopathy screening program; 96 images have discernable optic nerve head (Lowell et al., 2004). The subjects are from various ethnic backgrounds; 19 have type 2 diabetes mellitus, while the diabetes status was unavailable for the remaining 31. In this database there is considerable quality variation in the images, with many characteristics that can affect segmentation algorithms. The optic nerve head center is marked up a clinician, and four clinicians marked the optic nerve head edge where it intersects with radial spokes (at 15 angles) radiating from the nominated center.⁶

FOMDB

The Fundación Oftalmológica del Mediterráneo (Spain) provided us a private dataset of 67 fundus images (2048 x 1536) for hypertensive risk determination. The images belong to 67 subjects between 33 and 73 years old. Among them, a group without previous known pathology and a group of hypertensive patients were established. The first group was

⁵<http://messidor.crihan.fr/download-en.php>

⁶<http://reviewdb.lincoln.ac.uk/ImageDatasets/ONHSD.aspx>

formed by 38 healthy controls and the second group by 29 hypertensive patients previously diagnosed and treated or not for more than 5 years of evolution. Both groups were established based on the disease history of their members.

3.7.2 Retinal vessel segmentation

To validate the vessel segmentation, the results of the method of Subsection 3.3.1 were compared with the hand-segmented images from DRIVE database (Staal et al., 2004). The first-observer images were taken as reference (gold standard) to calculate similarity degree between them and our segmentation.

The performance of the method was evaluated based on three concepts: accuracy (Ac) and true positive (TPF) and false positive (FPF) fractions. Accuracy (Ac) is determined by the sum of correctly classified pixels as vessel and non-vessel divided by the total number of pixels in the image. True positive fraction (TPF), also known as sensitivity, is established by dividing the correctly classified pixels as vessel by the total number of vessel pixels in the gold standard. False positive fraction (FPF) is calculated by dividing the misclassified pixels as vessel by the total number of non-vessel pixels in the gold standard. In Table 3.1 these results can be observed. The images manually segmented by the second observer were also compared with the gold standard in order to note that there are differences between both experts. In summary, the average values obtained by the proposed method were: $Ac = 0.9417$, $TPF = 0.6570$ and $FPF = 0.0166$. Table 3.2 displays the true positive/negative and false positive/negative pixels used to calculate the measures of the Table 3.1.

Table 3.1: Comparison between the results of the method proposed in Subsection 3.3.1 and the 2nd observer of DRIVE database (average values and standard deviations) regarding the gold standard.

	Ac	TPF	FPF
Proposed method	0.9417 (0.0076)	0.6570 (0.0668)	0.0166 (0.0093)
2 nd observer	0.9473 (0.0048)	0.7757 (0.0596)	0.0275 (0.0083)

Table 3.2: True positives/negatives and false positives/negatives (TP, TN, FP and FN). Average values and standard deviations.

	TP	TN	FP	FN
Proposed method	18,945 (2,309)	194,740 (3,557)	3,284 (1,842)	9,938 (2,294)
2 nd observer	22,374 (2,389)	192,570 (3,083)	5,453 (1,685)	6,508 (1,895)

Figure 3.44 depicts some vessel segmentation results. The first two rows show good segmentation results where only the very thin vessels are not detected. In the last two rows, in addition to the non-detection of the very thin vessels, a bit of over-segmentation can be observed in some areas of the images.

On the other hand, in (Niemeijer et al., 2004) a study about the accuracy of different vessel segmentation methods was carried out on the same database. This study compared several approaches: matched filter (Chaudhuri et al., 1989), scale-space analysis and region growing (Martinez-Perez et al., 1999), mathematical morphology and curvature estimation (Zana and Klein, 2001), verification-based local thresholding (Jiang and Mojon, 2003) and pixel classification (Niemeijer et al., 2004). Table 3.3 shows a comparison between the aforementioned methods and ours. Our method achieved a higher accuracy ratio and its false positive fraction is, even, lower than the second observer fraction.

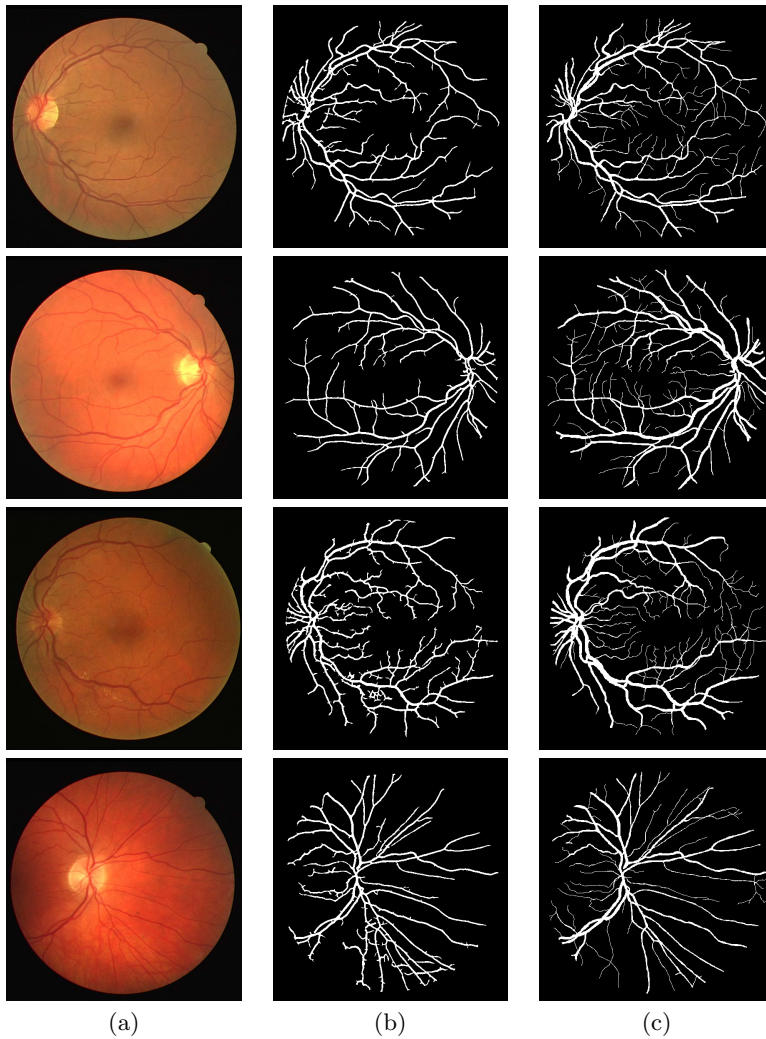


Figure 3.44: Some vessel segmentation results on DRIVE database: (a) Original fundus images, (b) Results of the method proposed in Subsection 3.3.1, and (c) Ground truth images.

Table 3.3: Accuracy (average and standard deviation) of several methods on DRIVE database.

	Ac
Proposed method	0.9417 (0.0076)
2 nd observer	0.9473 (0.0048)
Niemeijer <i>et al.</i> (Niemeijer <i>et al.</i> , 2004)	0.9416 (0.0065)
Zana and Klein (Zana and Klein, 2001)	0.9377 (0.0077)
Jiang and Mojón (Jiang and Mojon, 2003)	0.9212 (0.0076)
Martínez-Pérez <i>et al.</i> (Martinez-Perez <i>et al.</i> , 1999)	0.9181 (0.0240)
Chaudhuri <i>et al.</i> (Chaudhuri <i>et al.</i> , 1989)	0.8773 (0.0232)

In addition to improving the accuracy ratio, it must be stood out that the optic disk edge is not detected as vessel unlike the most edge detection methods. Figure 3.45 shows our method along with the proposed in (Martinez-Perez *et al.*, 2007). The non-detection of this feature is crucial to avoid multiple errors in any tracking process.

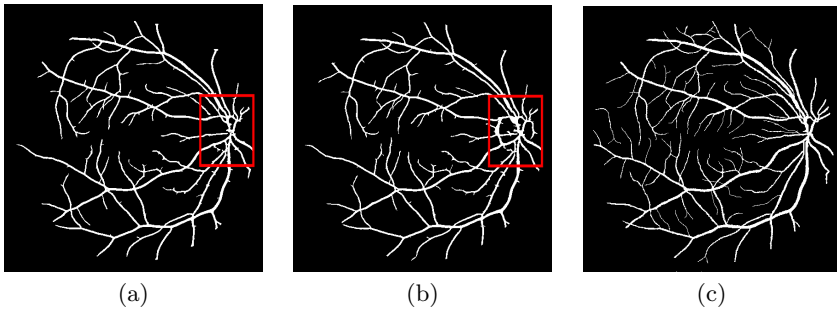


Figure 3.45: Vessel segmentation comparison: (a) Method proposed in Subsection 3.3.1, (b) Method proposed in (Martinez-Perez *et al.*, 2007) and (c) Manually segmented image belonging to the DRIVE database.

After the study performed by Niemeijer *et al.* (Niemeijer *et al.*, 2004), other methods about retinal vessel segmentation were published. As far the author knows, the best results on DRIVE database were presented by Mendonça and Campilho (Mendonça and Campilho, 2006)

and Ricci and Perfetti (Ricci and Perfetti, 2007). Although these works achieved a slightly higher accuracy than the proposed method, $Ac = 0.9463$ and $Ac = 0.9595$, respectively; the border of the optic disk was still detected as vessel.

To evaluate the effect of the input image in the segmentation method, different tests were performed. The use of the red and green component and PCA were compared, being the green component which obtained the best results as can be appreciated in Figure 3.46.

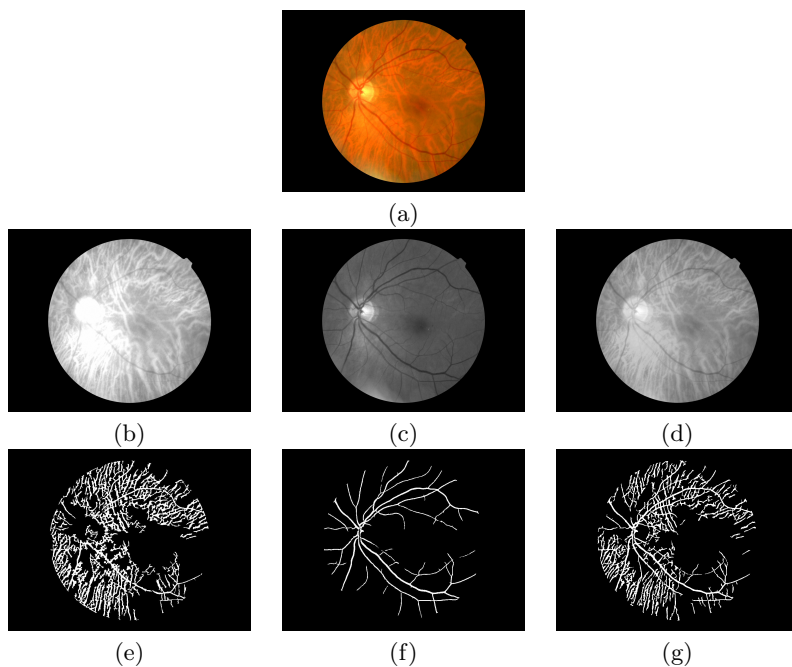


Figure 3.46: *Effect of the input image in the segmentation method: (a) Original RGB image, (b) Red component, (c) Green component, (d) First component of PCA, (e) Result of the method proposed in Subsection 3.3.1 on the red component, (f) Result of the method proposed in Subsection 3.3.1 on the green component and (g) Result of the method proposed in Subsection 3.3.1 on the first component of PCA.*

3.7.3 Retinal vessel centerline

The validation of the vessel centerline extraction method, that was presented in Subsection 3.3.2, was carried out on 2 public databases widely used: DRIVE (Staal et al., 2004) and STARE (Hoover et al., 2000). Although, in both databases, manual segmentations are included, the complete vasculature was detected, not only the vessel centerline which is the goal of our work. For that reason, the homotopic skeleton (Soille, 2003) associated to the manual segmentations was obtained for future comparisons. In Figure 3.47, the results of the proposed method on two images of DRIVE and STARE databases can be observed.

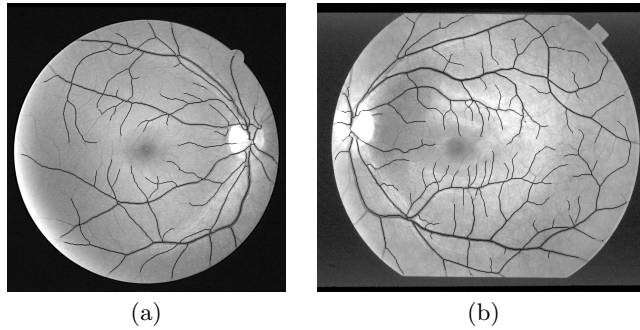


Figure 3.47: *Skeleton results of the method proposed in Subsection 3.3.2: (a) DRIVE image ('19_test') and (b) STARE image ('im0255').*

The validation was performed in two ways. One of them is based on comparing the results of this work with methods that first segment the vessels and after perform a skeletonization process and the other compares it with algorithms that directly obtain the skeleton.

On the one hand, regarding methods that require a previous segmentation, the presented algorithm was compared with two methods previously published. The first compared method uses the local maxima over scales of the magnitude of the gradient and the maximum principal curvature of the Hessian tensor in a multiple pass region growing

procedure (Martinez-Perez et al., 2007). The other compared method is that explained in Subsection 3.3.1. As mentioned before, both methods of the Subsection 3.3.1 and 3.3.2 are based on mathematical morphology and curvature evaluation although the morphological operations used are different as well as the obtained result. In the same way as explained above, the homotopic skeleton was performed after the segmentation process in both cases. On the other hand, as for the methods that obtain directly the retinal vessel centerline, the analysis was focused on other two approaches based also on the watershed transformation (Walter and Klein, 2002; Bessaid et al., 2009). In Figures 3.48 and 3.49, the strengths and weaknesses of the proposed method can be observed in two examples of both databases.

Avoiding complete vessel segmentation supposes an improvement in the automatic fundus processing since the skeleton is not dependent of a previous stage and the computational cost is reduced by decreasing the number of required steps. Apart from this fact, it must be stressed that an important advantage of the proposed method is its performance in pathological images or with large changes in illumination, as was observed in Figures 3.47, 3.48 and 3.49. In those cases, the presented algorithm works properly and reduces over-segmentation problems which can be found in methods based on a previous segmentation (Martinez-Perez et al., 2007; Morales et al., 2012). With regard to other methods that obtain the skeleton in a direct way and use the watershed transformation (Bessaid et al., 2009; Walter and Klein, 2002) instead of the stochastic watershed, the proposed work achieves a more robust detection and decreases the number of spurs. Despite good results, it must be mentioned that the main disadvantage of the method is that some vessels can lose their continuity if some part of them is not detected and it should be corrected in a post-processing stage.

3.7.4 Optic disk

The performance of the OD-segmentation method proposed in Section 3.5 was evaluated based on different concepts. Jaccard's

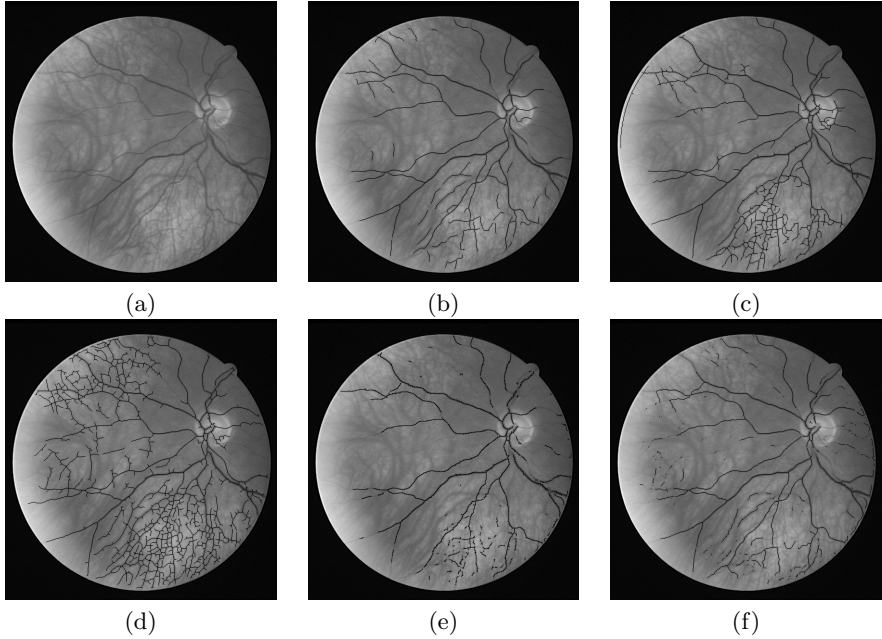


Figure 3.48: Comparison between different methods on DRIVE image ('23_training'): (a) Original image, (b) Method proposed in Subsection 3.3.2, (c) Martinez et al. method (Martinez-Perez et al., 2007), (d) Method presented in Subsection 3.3.1 (Morales et al., 2012), (e) Bessaid et al. method (Bessaid et al., 2009) and (f) Walter and Klein method (Walter and Klein, 2002).

(JC) and Dice's (S) coefficients describe similarity degree between two compared elements being equal to 1 when segmentation is perfect. Accuracy (Ac) is determined by the sum of correctly classified pixels as OD and non-OD divided by the total number of pixels in the image. True positive fraction (TPF) is established by dividing the correctly classified pixels as OD by the total number of OD pixels in the gold standard. False positive fraction (FPF) is calculated by dividing the misclassified pixels as OD by the total number of non-OD pixels in the gold standard. Finally, in order to be able to compare with more other authors' works, another measure was calculated: the mean absolute

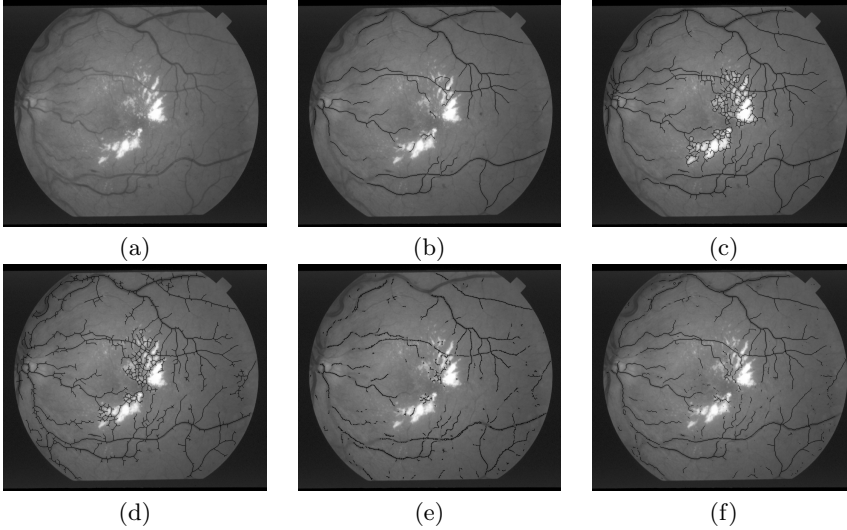


Figure 3.49: Comparison between different methods on STARE image ('im0001'): (a) Original image, (b) Method proposed in Subsection 3.3.2, (c) Martinez et al. method (Martinez-Perez et al., 2007), (d) Method presented in Subsection 3.3.1 (Morales et al., 2012), (e) Bessaid et al. method (Bessaid et al., 2009) and (f) Walter and Klein method (Walter and Klein, 2002).

distance (MAD) (Chalana et al., 1996), whose aim is to measure the accuracy of the OD-boundary,

$$MAD(\mathcal{A}, \mathcal{B}) = \frac{1}{2} \left\{ \frac{1}{n} \sum_{i=0}^n d(\mathbf{a}_i, \mathcal{B}) + \frac{1}{n} \sum_{i=0}^n d(\mathbf{b}_i, \mathcal{A}) \right\}, \quad (3.31)$$

where \mathcal{A} and \mathcal{B} are two curves corresponding to the border of the segmented OD and the border of the OD considered as ground truth, respectively. They are represented as set of points $\mathcal{A} = \{\mathbf{a}_1, \mathbf{a}_2, \dots, \mathbf{a}_n\}$ and $\mathcal{B} = \{\mathbf{b}_1, \mathbf{b}_2, \dots, \mathbf{b}_n\}$ and $d(\mathbf{a}_i, \mathcal{B})$ is the distance to the closest point for point \mathbf{a}_i on the curve \mathcal{B} .

In Table 3.4 the results on 5 public databases can be observed: DRIONS (Carmona et al., 2008), DIARETDB1 (Kauppi et al., 2007), DRIVE (Staal et al., 2004), MESSIDOR, and ONHSD (Lowell et al., 2004). Note that all these measures are indicators of the segmentation quality but in different way, i.e. a database can have a higher value of a measure than other database but this does not mean that this tendency must be followed by the rest. For example, the MAD can be equal in a case of under-segmentation and in other of over-segmentation if the distance between the boundary of our result and the ground truth (GT) is the same. As the OD is fitted by a circle, this situation may occur if our result and the ground truth were two concentric circles as shown in Figure 3.50. Although the MAD was equal, the rest of the measures could be different. Table 3.5 displays the true positive/negative and false positive/negative pixels used to calculate all measures of Table 3.4.

Table 3.4: Results (average values and standard deviations) obtained by the method proposed in Subsection 3.5 on DRIONS, DIARETDB1, DRIVE, MESSIDOR, and ONHSD databases. Jaccard's (JC) and Dice's (S) coefficients, accuracy (Ac), true positive (TPF) and true negative fractions (FPF), and mean absolute distance (MAD).

	JC	S	Ac
DRIONS	0.8424 (0.1174)	0.9084 (0.0982)	0.9934 (0.0051)
DIARETDB1	0.8173 (0.1308)	0.8930 (0.0913)	0.9957 (0.0039)
DRIVE	0.7163 (0.1880)	0.8169 (0.1712)	0.9903 (0.0134)
MESSIDOR	0.8228 (0.1384)	0.8950 (0.1056)	0.9949 (0.0050)
ONHSD	0.8045 (0.1175)	0.8867 (0.0776)	0.9941 (0.0042)
	TPF	FPF	MAD
DRIONS	0.9281 (0.1177)	0.0040 (0.0041)	4.1098 (3.4684)
DIARETDB1	0.9224 (0.1058)	0.0028 (0.0029)	9.6759 (8.4836)
DRIVE	0.8544 (0.1938)	0.0061 (0.0085)	7.9981 (9.9957)
MESSIDOR	0.9300 (0.1239)	0.0035 (0.0041)	5.8387 (6.5215)
ONHSD	0.9310 (0.1046)	0.0043 (0.0042)	4.4826 (3.0962)

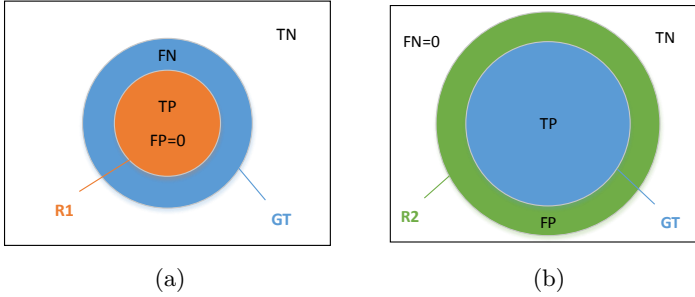


Figure 3.50: Sub and over-segmentation case: (a) Under-segmentation and (b) Over-segmentation. *GT* (in blue) is the ground truth in both cases and *R1* (in orange) and *R2* (in green) the result of our segmentation in each case. *R1* is characterized by not having false positives ($FP=0$) and *R2* by not having false negatives ($FN=0$). Moreover, the number of true positives (*TP*) will be greater in (b) than in (a) in contrast to the true negatives (*TN*). Although the *MAD* was equal for both cases, these variations could provoke that the rest of the measures were different.

Table 3.5: True positives/negatives and false positives/negatives (*TP*, *TN*, *FP* and *FN*). Average values and standard deviations.

	TP	TN	FP	FN
DRIONS	6,701 (1,392)	185,870 (3,467)	753 (763)	517 (877)
DIARETDB1	4,448 (875)	243,970 (1,087)	681 (705)	380 (542)
DRIVE	4,394 (1,252)	220,380 (2,867)	1,340 (1,869)	869 (1670)
MESSIDOR	9,773 (4,434)	427,450 (168,890)	1,415 (1,757)	804 (1,529)
ONHSD	4,667 (1,048)	196,240 (1,262)	846 (828)	345 (516)

On the one hand, in order to analyze the results on DRIONS database, the first observer images of this database were taken as reference (gold standard) to calculate similarity degree between them and our segmentation. Regarding MESSIDOR database, the OD rim of these 1200 images was hand segmented by the University of Huelva

(Spain) and it is currently available online to facilitate performance comparison between different methods.⁷ Concerning ONHSD dataset, the average of the edges marked by the four experts was used to generate the reference images. On the other hand, neither DIARETDB1 nor DRIVE databases have the OD segmented publicly available. In those cases, we compared our results with the same ground truth used in (Welfer et al., 2010), where the contour of each image was labelled by four ophthalmologists, and then, only the mean of those contours is considered as ground truth.

Apart from quantifying the accuracy in the OD boundary detection (Table 3.4), the accuracy in the disk localization was also measured. The performance of the optic disk localization is shown in Table 3.6. An automatically detected OD was considered correct if there exist intersection between it and the OD marked manually.

Table 3.6: *Success rate in the optic disk location.*

	Location performance (%)
DRIONS	100
DIARETDB1	100
DRIVE	97.50
MESSIDOR	95.83
ONHSD	100

Some examples of good, fair and bad OD segmentation results on the different databases analyzed are shown from Figure 3.51 to Figure 3.55).

⁷www.uhu.es/retinopathy/muestras.php

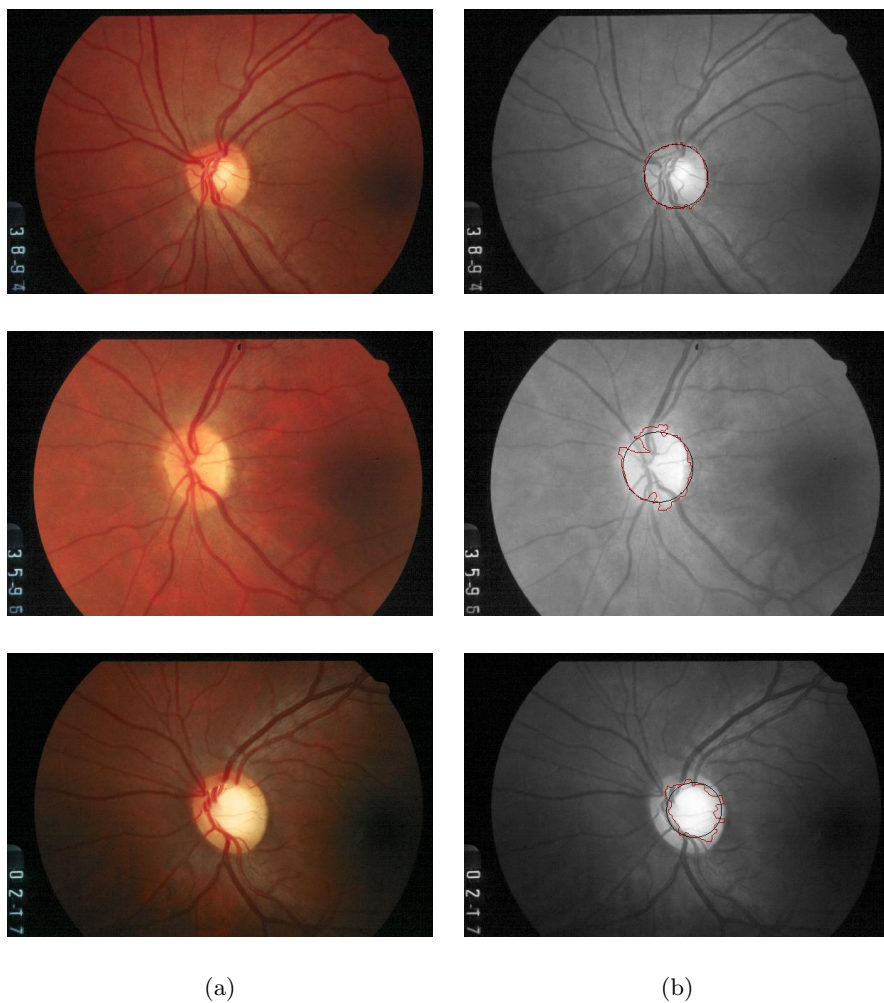


Figure 3.51: Some OD segmentation results on DRIONS database: (a) Original fundus images, (b) Results of the method proposed in Section 3.5 (OD contour in red and circular approximation of the OD contour in black). Top row: good segmentation. Middle row: fair segmentation. Bottom row: poor segmentation.

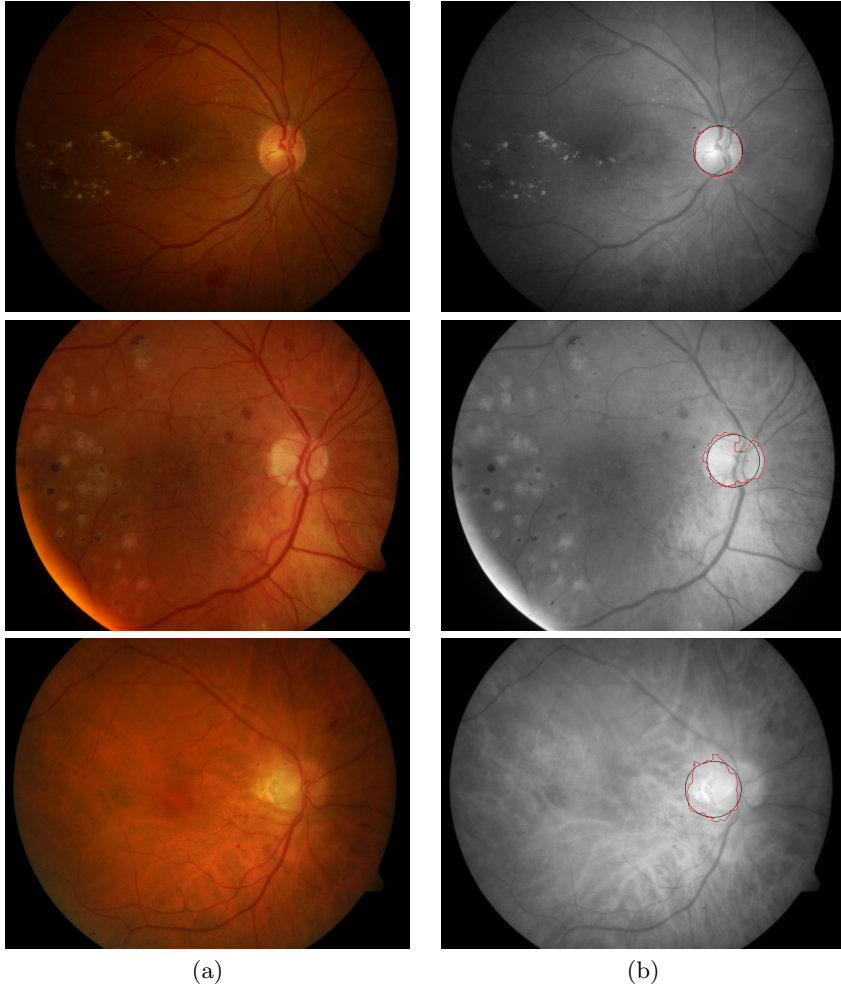


Figure 3.52: Some OD segmentation results on DIARETDB1 database: (a) Original fundus images, (b) Results of the method proposed in Section 3.5 (OD contour in red and circular approximation of the OD contour in black). Top row: good segmentation. Middle row: fair segmentation. Bottom row: poor segmentation.

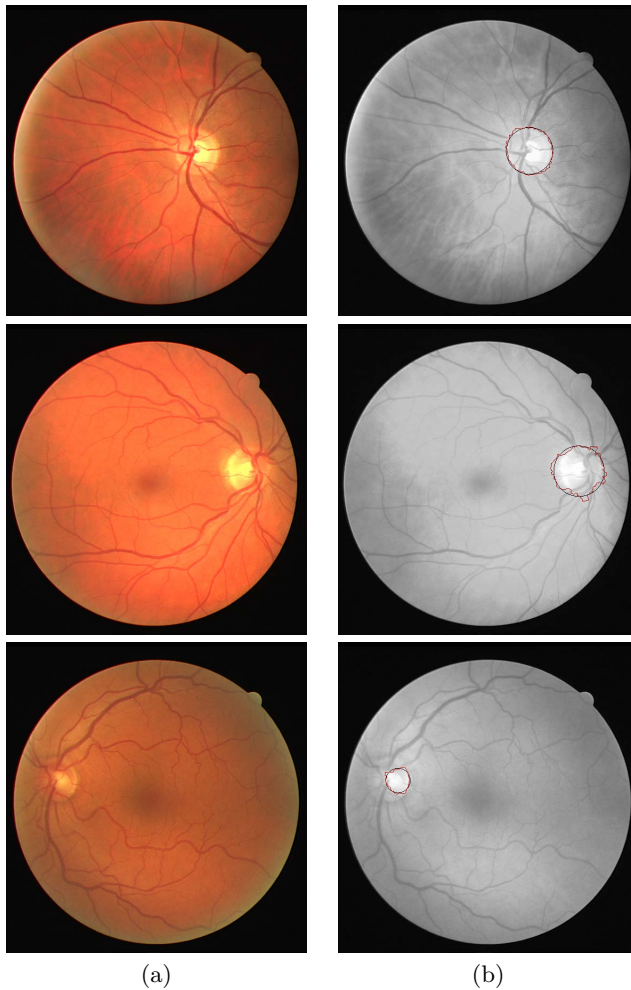


Figure 3.53: *Some OD segmentation results on DRIVE database: (a) Original fundus images, (b) Results of the method proposed in Section 3.5 (OD contour in red and circular approximation of the OD contour in black). Top row: good segmentation. Middle row: fair segmentation. Bottom row: poor segmentation.*

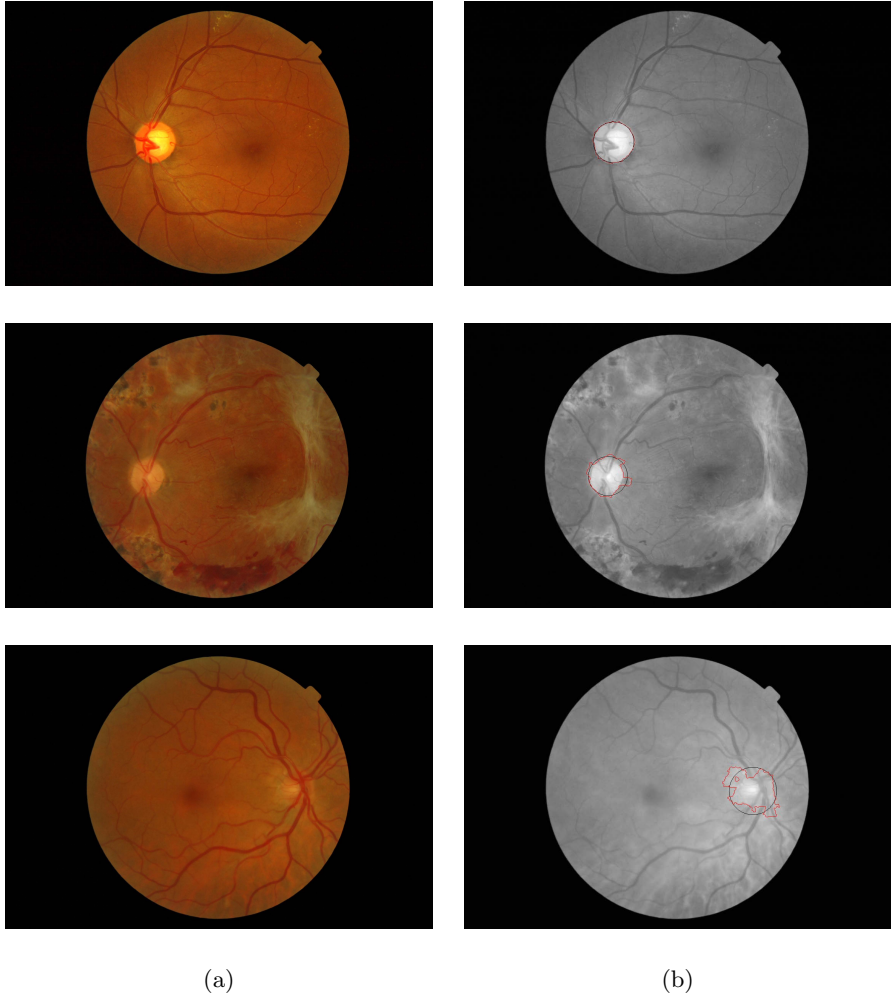


Figure 3.54: Some OD segmentation results on MESSIDOR database: (a) Original fundus images, (b) Results of the method proposed in Section 3.5 (OD contour in red and circular approximation of the OD contour in black). Top row: good segmentation. Middle row: fair segmentation. Bottom row: poor segmentation.

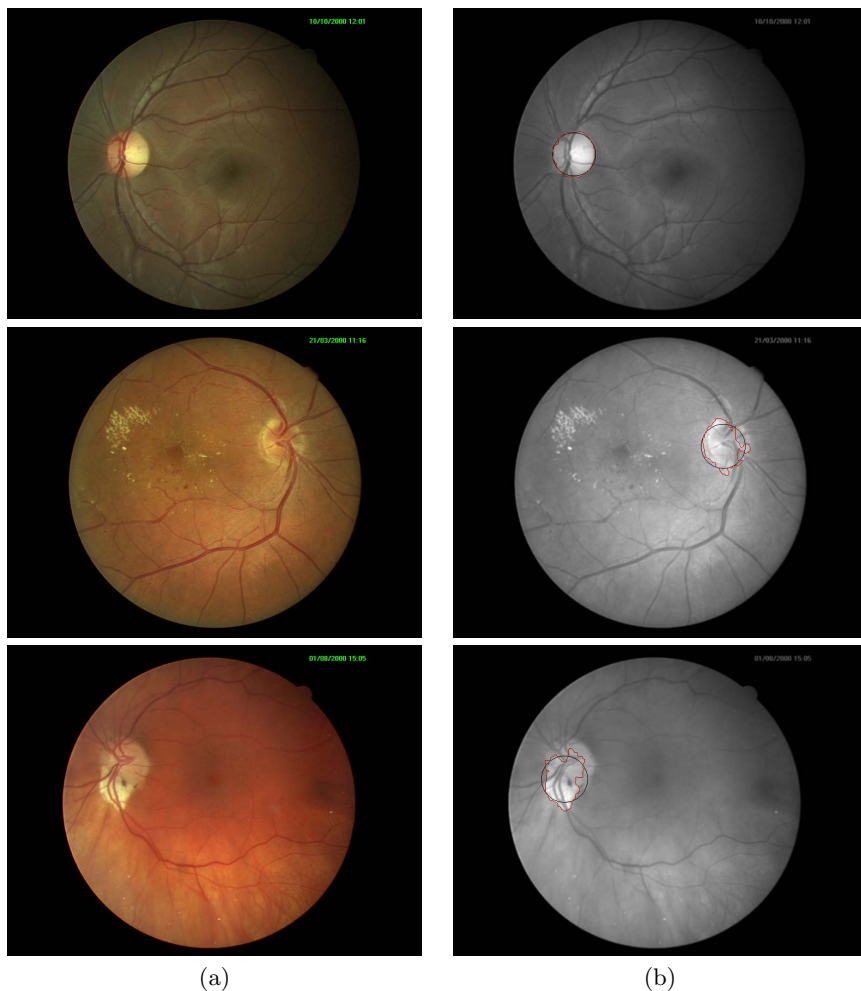


Figure 3.55: *Some OD segmentation results on ONHSD database: (a) Original fundus images, (b) Results of the method proposed in Section 3.5 (OD contour in red and circular approximation of the OD contour in black). Top row: good segmentation. Middle row: fair segmentation. Bottom row: poor segmentation.*

As was mentioned in Section 3.1, most OD segmentation methods are divided into morphological algorithms, template-based methods, and deformable models. The proposed algorithm was compared with several methods of each category. Most works were selected due to the fact they use some of the analyzed databases, and thereby, the current and future comparison is facilitated.

- *The presented method versus other morphological algorithms*

On the one hand, Table 3.7 analyses further DRIONS database at the same time that the performance of our work is contrasted with the performance of other method based on mathematical morphology. So, first, the segmented images by the second observer were also compared with the gold standard to obtain inter-expert differences, and secondly, other existing technique based on marker-controlled-watershed transformation (Walter et al., 2002) was implemented and compared with ours.

Table 3.7: Comparison of the parameters of Table 3.4 achieved by the method proposed in Section 3.5, by the 2nd observer and by other marker-controlled-watershed algorithm on DRIONS database.

	JC	S	Ac
Proposed method	0.8424 (0.1174)	0.9084 (0.0982)	0.9934 (0.0051)
2 nd observer	0.9202 (0.0455)	0.9578 (0.0265)	0.9970 (0.0017)
Walter <i>et al.</i> (Walter et al., 2002)	0.6227 (0.3695)	0.6813 (0.3854)	0.9689 (0.0492)
	TPF	FPF	MAD
Proposed method	0.9281 (0.1177)	0.0040 (0.0041)	4.1098 (3.4684)
2 nd observer	0.9498 (0.0537)	0.0012 (0.0009)	1.8887 (1.1455)
Walter <i>et al.</i> (Walter et al., 2002)	0.6715 (0.3980)	0.0210 (0.0417)	29.064 (48.058)

On the other hand, the concepts of Jaccard’s coefficient (also known as area overlap) and the MAD included in Table 3.4 allow us to compare with the method proposed in (Welfer et al., 2010) as well as with other state-of-the-art algorithms that were analyzed in it. Table 3.8 compares the results of the method presented in Section 3.5 with the method that achieves the best results in (Welfer et al., 2010) on the same databases (DIARETDB1 and DRIVE).

Table 3.8: Comparison of the method proposed in Section 3.5 with the work presented in (Welfer et al., 2010) based on Jaccard’s coefficient (JC) and mean absolute distance (MAD) on DIARETDB1 and DRIVE databases.

	DIARETDB1	
	JC	MAD
Proposed method	0.8173 (0.1308)	9.6759 (8.4836)
Welfer <i>et al.</i> (Welfer et al., 2010)	0.4365 (0.1091)	8.3100 (4.0500)
	DRIVE	
	JC	MAD
Proposed method	0.7163 (0.1880)	7.9981 (9.9957)
Welfer <i>et al.</i> (Welfer et al., 2010)	0.4147 (0.0833)	5.7400

According to these data, the method of the Section 3.5 obtains a mean overlap greater than 70% for the five databases which significantly improves the results of the compared methods. As for MAD values, the results of Welfer *et al.* are slightly better. As the ground truth images of DIARETDB1 were resized from 1500 x 1152 pixels to 640 x 480, our results were also resized to have the same spatial resolution of ground truth and make the comparison possible.

- *The presented method versus template-based methods*

The validation on MESSIDOR database was performed through the comparison with a circular template-based method and four elliptical

template-based approaches (Aquino et al., 2010). The methods analyzed in this work can be compared in based on Jaccard’s coefficient. Three of the elliptical template-based variants were based on minimizing the algebraic distance (achieving a $JC \simeq 0.66$), and one was based on minimizing the geometric distance ($JC = 0.67$). Our method improves the performance of all elliptical approaches and achieves comparable results with the circular template-based method ($JC = 0.86$).

- *The presented method versus deformable models*

Our method was also tested on ONHSD database in order to compare with the tree approaches based on deformable models evaluated in (Lowell et al., 2004). In addition, with the aim of achieving a rigorous comparison, the same metric used in this work was calculated. A subjective perception of quality, which is based on four categories, was defined as a way of classifying the performance. For that, they defined the discrepancy as $\delta^j = \sum_i \frac{|m_i^j - \mu_i^j|}{\sigma_i^j + \epsilon}$, where μ_i^j and σ_i^j summarize the clinician’s choice of rim location on spoke i of image j . Division by σ compensates for uncertainty in rim position; $\epsilon = 0.5$ is a small factor to prevent division by zero where the clinicians are in exact agreement. Each image is classified as *Excellent*, *Good*, *Fair*, or *Poor* depending on the discrepancy value (up to one, two, five, or more, respectively).

Table 3.9 summarizes the performance of our method on the subjective classification of the ONHSD dataset so that it is directly comparable with the results presented in (Lowell et al., 2004). While the best approach of this work has *Excellent-Fair* performance in 83% of cases, our method obtains it in 94%.

Table 3.9: *Subjective classification of performance on ONHSD dataset.*

	Excellent	Good	Fair	Poor	Excellent-Fair
Proposed method	28%	36%	31%	6%	94%
TemploralLock (Lowell et al., 2004)	42%	31%	10%	17%	83%
Simple (Lowell et al., 2004)	9%	8%	30%	53%	47%
DV-Hough (Lowell et al., 2004)	39%	22%	20%	19%	81%

In addition to the comparison with other methods, different experiments were performed to demonstrate the robustness of the proposed algorithm:

1. The variability between fundus images in color, intensity, size, the presence of artefacts, etc. makes each state-of-the-art method uses a different input image: green (Lalonde et al., 2001; Eswaran et al., 2008; Niemeijer et al., 2009) and red (Walter et al., 2002; Hajer et al., 2008; Welfer et al., 2010) band of the original RGB image, or even a combination of both of them (Aquino et al., 2010; Lu, 2011), intensity component extracted from the HSI representation (Lowell et al., 2004) and lightness channel of the HLS space (Osareh et al., 2002). However, due to this fundus image variability, they do not always provided the desired results. Therefore a PCA, able to maximize the separation between the different objects of the image, was proposed as a more appropriate input image. For example, in Figure 3.56, PCA is compared with the use of the red component on a specific image. It can be observed that while the red component is completely over-saturated, PCA obtains a gray image where the OD could be segmented. Specifically, in Table 3.10 a quantitative analysis about the use of those two color planes can be observed concluding that the employment of PCA improves the final results.

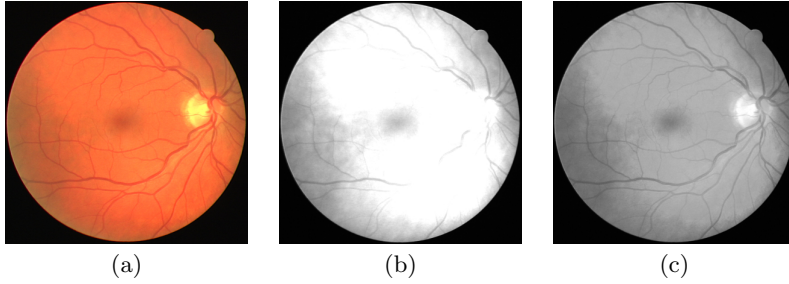


Figure 3.56: Advantages of PCA: (a) Original RGB fundus image, (b) Red component and (c) Image obtained by PCA.

Table 3.10: Comparison between the use of PCA versus the use of the red component on DRIVE database.

	PCA	Red Component
JC	0.7163 (0.1880)	0.4106 (0.3926)
S	0.8169 (0.1712)	0.4674 (0.4256)
Ac	0.9903 (0.0134)	0.8979 (0.2044)
TPF	0.8544 (0.1938)	0.5607 (0.4368)
FPF	0.0061 (0.0085)	0.0994 (0.2343)
MAD	7.9981 (9.9957)	72.741 (87.216)

- As for the improvements achieved by the proposed method in relation to others which use watershed transformation must also be highlighted. In particular, the different steps of our method help to avoid under-segmentation problems, as occurs in Figure 3.57 where the original image was segmented using only one internal marker located in the geodesic center of its largest and brightest object (Walter et al., 2002). It can be seen that there exist under-segmentation although the vessels were removed before applying the watershed transformation. Two different methods were used for vessel removal: a morphological closing and the inpainting technique presented in Subsection 3.5.1.

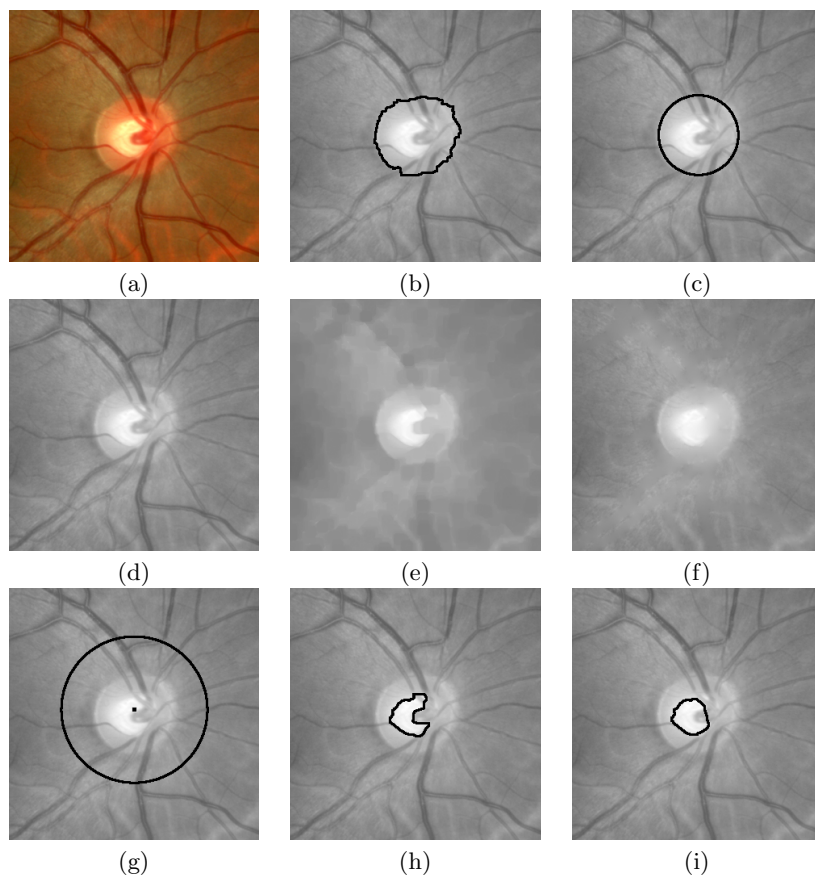


Figure 3.57: *Under-segmentation problem produced in other marker-controlled-watershed method: (a) Original RGB image, (b) Watershed result of the method proposed in Section 3.5, (c) Circular approximation of (b), (d) Grayscale image, (e) Vessel removal through a morphological closing, (f) Vessel removal through inpainting technique, (g) Internal and external marker, (h) Watershed region obtained on the gradient of (e) using the markers shown in (g) and (i) Watershed region obtained on the gradient of (f) using the markers shown in (g).*

3. Referring to the databases employed, most of the images included in them are pathological and therefore can distort the OD detection. For example, the images of patients with diabetic retinopathy may contain exudates which are of similar intensity to the OD or patients with glaucoma can suffer from peripapillary atrophy. Concretely, Figure 3.58 shows the robustness of the method in presence of exudates. However, if the lesion is as big as the optic disk and brighter, it is true that the method may fail.

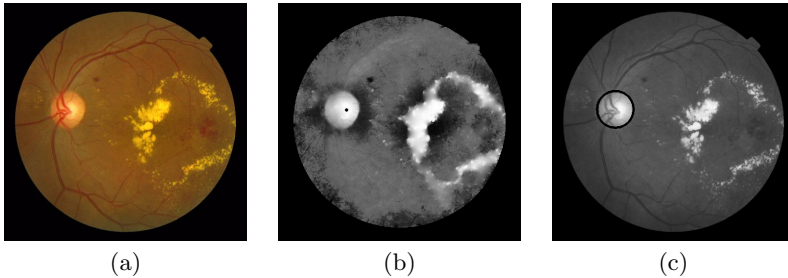


Figure 3.58: *Robustness of the gray-image centroid in presence of exudates: (a) Original image, (b) First PC (z_1) after inpainting technique with the centroid in black, and (d) Result of the proposed method.*

3.7.5 Computer-aided diagnosis software for hypertensive risk determination

For the system validation presented in Section 3.6, a set of 67 fundus images was used. These images belong to a private database of the Fundación Oftalmológica del Mediterráneo (FOMDB). The implemented system was installed in this organization in order to be used by its clinicians in their daily practice and be able to carry out a clinic validation of the developed software based on a double-blind study. The fundus images were evaluated twice. Once, by qualified ophthalmologists who determined the presence or absence of vascular alteration through visual inspection of the original image, and another time using the developed system. A patient was considered hypertensive

if the clinician detected in their fundus at least one sign of pathological arteriovenous crossing. The same three ophthalmologists rated the entire dataset with and without the software. A minimum period of one month was established between both rates to avoid the effect of the repeated exposure to the images. The final expert rates were obtained taking the majority vote among the three clinicians.

With the implemented tool, bifurcation angles, inside of an existing region of the original image concentric to the optic disk, were measured along with the caliber of a vein and an artery manually selected and situated at the same distance from the optic disk. Subsequently, a statistical analysis on different variables, extracted from data provided by the system, was conducted to see if any of them were able to discriminate whether a patient belonged to the group of hypertensive patients or to subjects without cardiovascular disease. Statistical Package for the Social Sciences (SPSS, IBM SPSS Data Collection) version 17.0 was used for this purpose.

Two parallel statistical studies were performed, one analyzing the variables related to the caliber and other analyzing those related to the bifurcation angles. The reason for the distinction was that only fundus images with at least five measured bifurcation angles were considered valid for this specific study. In both, the comparison between control and study groups was conducted using one-way ANOVA, where the dependent variables were each of the parameters calculated from the data provided by the software (vein caliber, artery caliber, relative caliber of the vein and artery (A/V ratio), average, median, deviation and variance of the branching angles, minimum and maximum angle and difference between the maximum and minimum angle) and the factor was the group formed by the subgroup without known disease and the subgroup of hypertensive patients.

Previous to the analysis, the normality of the calculated parameters was checked using the Kolmogorov-Smirnov test. From the obtained values, it can be concluded that only the caliber of the artery

($F(1;66) = 4.471; p < 0.05$) and the ratio between the caliber of the vein and the artery ($F(1;66) = 4.161; p < 0.05$) show statistically significant differences between both subgroups. Afterwards, using only these parameters, the optimal threshold to separate both classes was established and then the sensitivity, specificity and accuracy of the system to discriminate between healthy subjects and patients and with hypertensive pathology were calculated. Sensitivity and specificity measures the proportion of positives and negatives cases (pathological and healthy) which are correctly identified, respectively ($Sens = 56.41\%$ and $Spec = 67.86\%$). Accuracy is the proportion of true results (both true positives and true negatives) among the total number of cases examined ($Ac = 61.19\%$).

If the proportion of true results of the analyzed population is taken into account (i.e. the system accuracy), ophthalmologist visual diagnoses agreed with patient history 42% of the time without the computer assistance and that value increases up to 61% with the computer assistance. Their sensitivity and specificity are also improved from 38% and 50% to 56% and 68%, respectively. Therefore, it can be concluded that the use of this computer-aided diagnosis software provides an improvement of almost 20% in hypertension detection.

Thanks to the software validation, it was demonstrated that the caliber of the arteries and the relative caliber of the veins and arteries show significant differences when patients are classified either healthy or hypertensive. Despite these results, it cannot be concluded that bifurcation angles are not significant for this purpose because only 54% of the fundus of the dataset could be analyzed due to the fact that the remainder contains less than 5 angles per image. Thus, the region where the measures are taken, should be enlarged or the database increased to repeat the same analysis.

Although the values of accuracy, sensitivity and specificity of the system are not too much high in the hypertension discrimination, it must be stressed that the results achieved by clinicians by visual

inspection of the fundus are improved almost 20%.

With regard to the obtained measures, they are accurate and reliable but also dependent on a correct skeleton detection and significant point classification.

3.8 Conclusions

In this chapter, several methods related to the detection and characterization of the retinal vessels and optic disk from a fundus image have been presented.

The method proposed in Subsection 3.3.1 for vessel segmentation is based mainly on morphological operators but also used principal curvature information, allowing an enhanced detection of the vessels on to the background of the image. This algorithm was compared with several methods and, apart from being among the best of the state of the art ($Ac = 0.9417$), the optic disk edge is not detected as vessel, which facilitates a later vessel tracking process.

Subsection 3.3.2 deals with a method for vascular skeleton determination. This work proposes a new approach based on mathematical morphology and curvature evaluation and makes use of the stochastic watershed to extract the vessel centerline in a direct way. A correct vessel skeleton detection is usually required to analyze different vessel features that can indicate the presence of several diseases. Avoiding complete vessel segmentation supposes an improvement since the skeleton is not dependent of a previous stage at the same time that works properly in pathological images or with large changes in illumination and reduces over-segmentation problems of state-of-the-art methods based on a previous segmentation. With regard to other methods that obtain the skeleton in a direct way and use the marker-controlled watershed transformation, the stochastic watershed achieves a more robust detection and decreases the number of spurs.

In Section 3.5, a new approach for the automatic detection of the optic disk was introduced. First, it is focused on the use of a new gray image as input obtained through PCA which combines the most significant information of the three RGB components. Secondly, several operations based on mathematical morphology are implemented with the aim of locating the OD. For that purpose, both stochastic and stratified watershed as well as geodesic transformations, such as gray level distance and gray level close holes, were used. The algorithm was validated on 5 different public databases obtaining promising results and improving the results of other methods of the literature ($JC = 0.8200$, $S = 0.8932$, $Ac = 0.9947$, $TPF = 0.9275$, $FPF = 0.0036$, and $MAD = 5.9060$). These values correspond with the weighted average of the results of the proposed method, i.e., they are calculated according to the number of images of each analyzed database.

Making use of the aforementioned methods, a computer-aided diagnosis software for hypertensive risk determination was presented in Section 3.6. From a fundus image, the tool automatically detects blood vessels and allows measurement of bifurcation angles and selection of branches to determine their caliber. These data facilitate expert medical diagnosis and study of the progression of the disease. In particular, the software was used by expert ophthalmologists to help them to discriminate between a normal vascularization and cardiovascular pathology. The accuracy, sensitivity and specificity of the hypertension detection achieved by direct visual inspection of the fundus were improved by almost 20%.

Chapter 4

Retina background characterization

This chapter focuses on analyzing the texture of the fundus images by means of local binary patterns to differentiate between pathological and healthy images. In particular, its goal is to distinguish between diabetic retinopathy (DR), age-related macular degeneration (AMD) and normal fundus images analyzing directly the retina background texture and avoiding a previous lesion segmentation stage. Chapter 4 arose from a research stay in the University of Stavanger under the supervision of Prof. Kjersti Engan.

The information contained in this chapter was included in a journal paper that is currently under revision:

Morales, S., Engan, K., Naranjo, V., Colomer, A. (2015). Retinal Disease Screening through Local Binary Patterns. *IEEE Journal of Biomedical and Health Informatics*.

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4.1 Introduction

Diabetic retinopathy (DR) and age-related macular degeneration (AMD) are nowadays two of the most frequent causes of blindness and vision loss (World Health Organization, 2010a). In addition, these diseases will experience a high growth in the future due to diabetes incidence increase and aging population in the current society. Based on these facts, a computer-aided diagnosis software capable of discriminating, through image processing, between a healthy fundus (without any pathology) and DR and AMD patients is presented in this chapter. Specifically, the aim of the proposed software is to carry out an automatic screening of these diseases facilitating the at-risk population assessment.

DR and AMD can be characterized by the presence of specific types of retinal lesions such as microaneurysms, exudates or drusen, among others, as was seen in Section 2.2. Figure 4.1 depicts some examples of these diseases in comparison with the fundus image from a healthy subject.

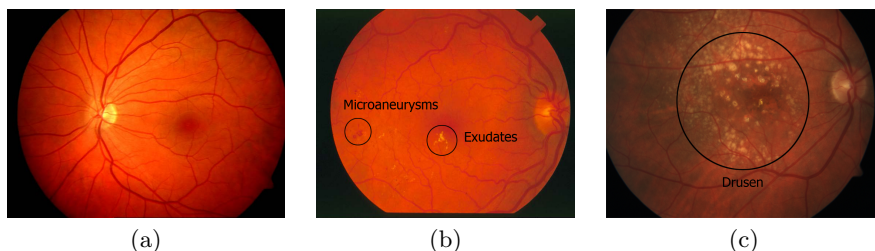


Figure 4.1: *Fundus images. (a) Healthy, (b) DR (with microaneurysms and exudates) and (c) AMD (with drusen).*

This chapter focuses on investigating the discrimination capabilities in the fundus texture to differentiate between pathological and healthy images. In particular, the main focus lies in exploring the performance of Local Binary Patterns (LBP) as a texture descriptor for retinal

images. LBP is a technique used for describing textures that has been given a lot of attention in recent years (Ojala *et al.*, 2001; Ojala *et al.*, 2002). It is based on looking at the local variations around each pixel, and assigning labels to different local patterns. Thereafter, the distribution of the labels is evaluated and used in the classification stage. There are many examples of the success of LBP used to describe and classify textures in general (Ahonen *et al.*, 2006; Heikkil 2009; Liao *et al.*, 2009; Yang and Ai, 2007) and also in the case of medical imaging (Kotu *et al.*, 2012; Oppedal *et al.*, 2012; Nanni *et al.*, 2010). However, regarding fundus image processing, LBP has not been widely used. Most state-of-the-art works that use the LBP technique on fundus images focus on the segmentation of the retinal vessels (Zabihi *et al.*, 2010; Dhanushkodi and Vasuki, 2013) rather than on a full diagnosis system, although some examples can be found in this direction (Garnier *et al.*, 2014; Mookiah *et al.*, 2013). Garnier *et al.* deal with the AMD detection using LBP. The texture information on several scales is analyzed through wavelet decomposition and the LBP histogram is found from the wavelet coefficients. Linear Discriminant Analysis (LDA) is used for feature dimension reduction using the values of the entire LBP histogram as input features. Image classification on a set of 45 images is evaluated with a leave-one-out validation method (Garnier *et al.*, 2014). Mookiah *et al.* extract abnormal signs from fundus images to detect normal fundus and two DR stages. Thirteen features, such as the area of hard exudates, the area of blood vessels, bifurcation points, textures and entropies, fed three different classifiers (Probabilistic Neural Network (PNN), Decision Tree C4.5 and Support Vector Machine (SVM)). The textures are found by LBP and Laws energy. A previous segmentation of the exudates, optic disk and blood vessels is needed for feature extraction. The experiments were conducted on 156 subjects and the PNN was chosen as the best classifier with three-fold cross validation (Mookiah *et al.*, 2013).

The goal of this chapter is to distinguish between DR, AMD and normal fundus images at the same time and avoiding any previous segmentation stage of retinal lesions. The texture of the retina

background is directly analyzed by means of LBP, and only this information is used to differentiate healthy patients and these two pathologies. A comprehensive study about what type of classifier obtains the best results is also undertaken. The performance of Logistic Regression, Neural Networks, SVM, Naive Bayes, J48, Rotation Forest, Random Forest and AdaBoost M1 is compared. This approach is different from previous works. Mookiah *et al.* require the segmentation of exudates in addition to segmentation of main structures (optic disk and vessels) for feature extraction and, although three different classes are identified, they only focus on DR detection. Garnier *et al.* do not need previous segmentations but only handle with AMD diagnosis. Both works use directly the information of the LBP histograms. However, in this work, LBP values are combined with a contrast measure and, instead of using directly the histograms values, different statistics are calculated from them.

The rest of the chapter is organized as follows: in Section 4.2 theoretical background is included and in Section 4.3 the proposed method is presented. Section 4.4 shows how system validation was performed and as well as the obtained results. Finally, Section 4.5 provides conclusions and some future areas for work.

4.2 Theoretical background

As Section 3.2, this section is intended as a guide to the reader about the main tool used in the chapter: local binary pattern theory.

Local binary pattern (LBP) is a powerful gray-scale texture operator used in many computer vision applications because of its computation simplicity (Ojala *et al.*, 2001; Ojala *et al.*, 2002). The first step in LBP is to produce a label for each pixel in the image where the label is found based on the local neighborhood of the pixel which is defined by a radius, R , and a number of points, P . The neighboring pixels are thresholded with respect to the gray value of the central pixel of the neighborhood generating a binary string or, in other words, a binary pattern. The

value of a LBP label is obtained for every pixel by summing the binary string weighted with powers of two. It can be represented as follows:

$$LBP_{P,R} = \sum_{p=0}^{P-1} s(g_p - g_c) \cdot 2^p, \quad s(x) = \begin{cases} 1 & \text{if } x \geq 0 \\ 0 & \text{if } x < 0. \end{cases} \quad (4.1)$$

where g_p and g_c are the gray values of the neighborhood and central pixel, respectively. P represents the number of samples on the symmetric circular neighborhood of radius R . The g_p values are interpolated to fit with a given R and P . The values of the labels depend on the size of the neighborhood (P). In Figure 4.2 two neighborhoods with different radii, but the same number of samples, are drawn.

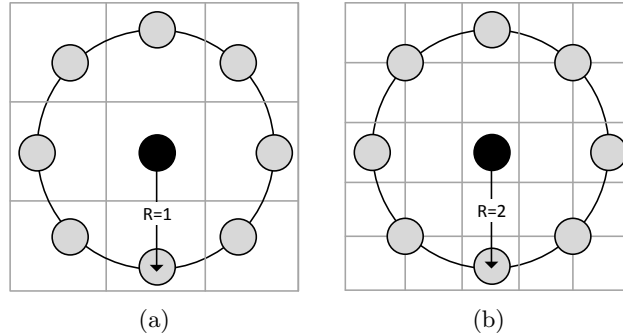


Figure 4.2: LBP neighborhoods: (a) $R = 1$ and $P = 8$. (b) $R = 2$ and $P = 8$.

2^P different binary patterns can be generated in each neighborhood. However, the bits of these patterns must be rotated to the minimum value to achieve a rotation invariant pattern. For example, the binary patterns 01110000, 10000011 and 00111000 are different rotations of the same pattern and correspond to the normalized string 00000111. In the case of $P = 8$, only 36 of the 2^P possible patterns are rotation invariant, i.e., $LBP_{8,R}$ can have 36 different values. Figure 4.3 shows

how LBP is calculated for a circular neighborhood of radius 1 ($R = 1$) and 8 samples ($P = 8$).

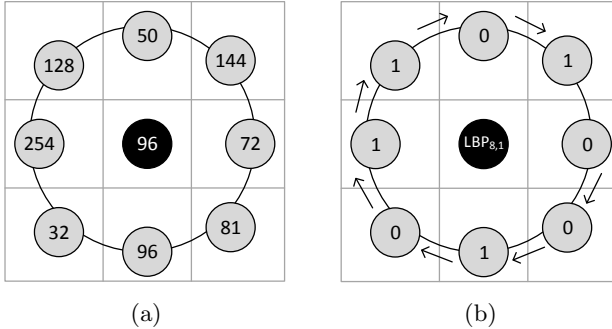


Figure 4.3: LBP computation: (a) Gray values of a circular neighborhood of radius 1 and 8 samples. (b) Thresholding between the gray value of the neighborhood and the central pixel. The rotation invariant local binary pattern generated is 00101101 (the arrows indicate the order in which the string is formed). Specifically, the LBP label is obtained as follows $LBP_{8,1} = 0 \times 2^0 + 0 \times 2^1 + 1 \times 2^2 + 0 \times 2^3 + 1 \times 2^4 + 1 \times 2^5 + 0 \times 2^6 + 1 \times 2^7 = 180$.

When LBP is used for texture description, it is common to include a contrast measure by defining the rotational invariant local variance as follows:

$$VAR_{P,R} = \frac{1}{P} \sum_{p=0}^{P-1} (g_p - \mu)^2, \quad \mu = \frac{1}{P} \sum_{p=0}^{P-1} g_p. \quad (4.2)$$

The LBP and VAR measures are complementary and are combined to enhance the performance of the LBP operator. The implementation of both measures is publicly available online.¹

¹www.ee.oulu.fi/~gyzhao/LBP_Book.htm

4.3 Application to pathology detection: Diabetic retinopathy and age-related macular degeneration

Using the technique explained above, an algorithm able to distinguish between healthy retinal images and images belonging to patients with diabetic retinopathy and age-related macular degeneration was developed.

4.3.1 General description

The algorithm proposed for classification of retina images does not need a prior segmentation of suspicious lesions. Manual segmentation of lesions are time consuming and automatic segmentation algorithms might not be accurate, thus removing the need for lesion segmentation can make the classification more robust. The algorithm is based on the study of the retina background. Hence, the main structures of the fundus (the vascular network and the optic disk), which are not related to the diseases under study, should not be taken into account when the texture of the fundus is analyzed. If these predominant structures were included in the image texture analysis, they could mask the difference between healthy and pathological images because their aspect in both cases would be similar.

4.3.2 Feature extraction

The LBP and VAR operators described above are used to characterize the texture of the retina background. They are calculated for each pixel of the RGB images using $P = 8$ and different values of R ($R = \{1, 2, 3, 5\}$). Due to the fact that the images under study belong to different databases, the size of the images varies. As the LBP and VAR depend on the radius of the neighborhood, the images must be resized to a standardized size to obtain comparable texture descriptors. The images are resized using the length of the horizontal diameter of

the fundus as reference (Zhang et al., 2012). Bicubic interpolation is used for resizing; the output pixel value is a weighted average of pixels in the nearest 4-by-4 neighborhood.

The red, green and blue components of each image are independently analyzed. One example of the aspect of the LBP and VAR images of an AMD fundus is represented in Figure 4.4. Only the pixels of the retina background are considered significant for the texture analysis. Thus the LBP and VAR values corresponding to pixel positions of the vessels, the optic disk or outside the fundus should not be taken into account. The vascular network and the optic disk are detected by the methods presented in Subsection 3.3.1 and Section 3.5 and, subsequently, the LBP and VAR labels are removed according to the pixel position of the optic disk and the vascular network. The external mask is directly obtained by thresholding.

The resulting LBP and VAR images provide a description of the image texture. After masking the optic disk and vessel segments, the LBP and VAR values within the external mask of the fundus are collected into histograms, one for each color (RGB). Different statistical information is extracted from these histograms to use it as features in the classification stage. Concretely, the calculated statistical values are: mean, standard deviation, median, entropy, skewness and kurtosis. To sum up, 6 statistical values are calculated from each LBP and VAR histogram, giving place to 12 features for each radius used. Consequently, the total number of features is equal to 144 (12 features x 4 radius x 3 components). Figure 4.5 depicts the feature extraction flowchart and Algorithm 4.1 summarizes the steps to be followed.

4.3.3 Classification

The dataset analyzed was divided into two subsets, one for training and testing by cross validation (model set) and other purely for testing (validation set).

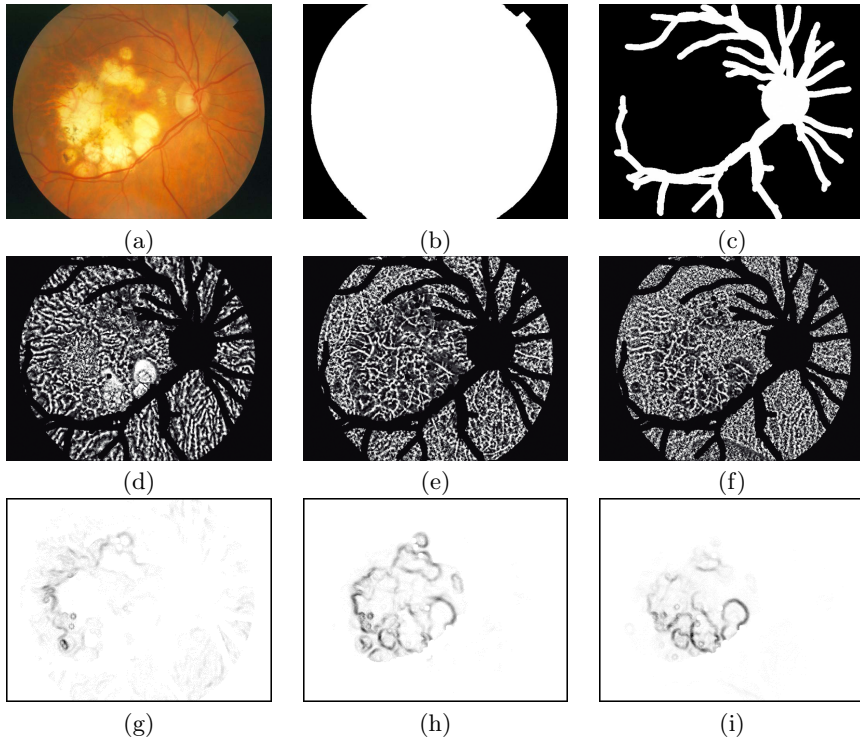


Figure 4.4: Feature extraction using $P = 8$ and $R = 5$. (a) AMD fundus image, (b) External mask, (c) Optic disk and vessel mask, (d-f) LBP images calculated on R , G and B components, respectively. Optic disk and vessel segments are removed (black). (g-i) VAR images calculated on R , G and B components, respectively. Optic disk and vessel segments are removed (white).

Once the features are extracted, the data of the model set must be preprocessed before the classification stage. In the preprocessing, two tasks are carried out: data normalization and data resampling. The first one because the range of values of raw data varies widely and the second one because the dataset is clearly unbalanced and most machine learning algorithms would not work properly. In particular, the method used for the normalization is to standardize all numeric attributes in the given

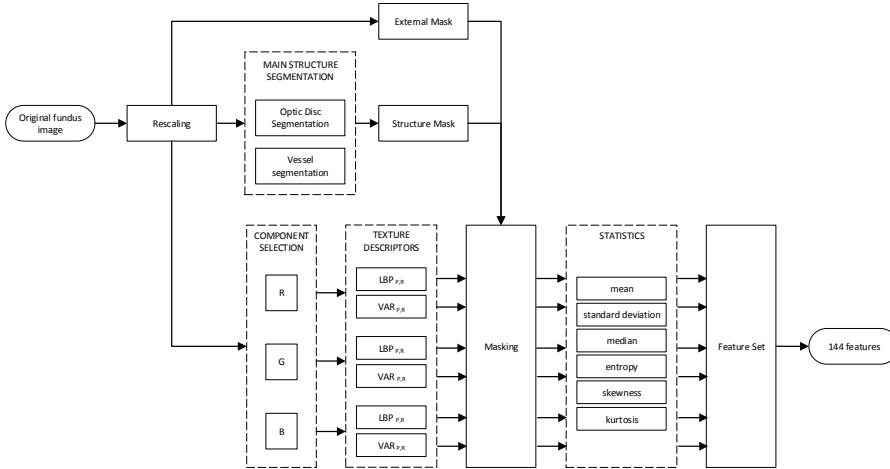


Figure 4.5: Feature extraction flowchart. First, the original fundus image is rescaled based on its horizontal diameter. Secondly, LBP and VAR measures are performed on the three RGB components using $P = 8$ and $R = \{1, 2, 3, 5\}$. Then, the external mask of the fundus is determined and the optic disk and vessels are segmented generating a structure mask. Both, external and structure masks, are used for masking the result of the texture descriptors. The final feature set is formed by the statistical values of all texture descriptors after masking.

dataset to have zero mean and unit variance and, for the resampling, the Synthetic Minority Oversampling TEchnique (SMOTE) (Chawla et al., 2002) is applied. This technique over-samples the minority class creating synthetic minority class examples to balance the dataset.

Afterwards, external cross validation (CV), also called nested CV (Scheffer, 1999; Dudoit and van der Laan, 2005), is performed on the model set so that the dimensionality of the data is reduced by feature selection before being passed on to a classifier. 10 folds are used in the external loop and 5 folds in the internal loop. The purpose of the internal loop is to select a feature subset and the used technique is

Algorithm 4.1: Feature extraction

Data: Original RGB fundus image $\mathbf{f}(f_r, f_g, f_b)$, External mask M_{ext} ,
Optic disk mask M_{OD} , Vessel mask M_v

Result: Feature vector, \mathbf{F}

initialization: P, \mathbf{R} ;

rescaling:

$\mathbf{f}'(f'_r, f'_g, f'_b) \leftarrow \text{resize}(\mathbf{f}(f_r, f_g, f_b))$;

$M'_{ext} \leftarrow \text{resize}(M_{ext})$;

$M'_{OD} \leftarrow \text{resize}(M_{OD})$;

$M'_v \leftarrow \text{resize}(M_v)$;

mask computing:

$M \leftarrow \text{and}(M'_{ext})^c, (M'_{OD})^c, (M'_v)^c$;

texture descriptors:

for $i = \{r, g, b\}$ **do**

for $j \leftarrow 1$ **to** $\text{length}(\mathbf{R})$ **do**

$f_{LBP_{ij}} \leftarrow LBP_{P, R_j}(f'_i)$;

$f_{VAR_{ij}} \leftarrow VAR_{P, R_j}(f'_i)$;

 statistics:

$F_{ij1} \leftarrow \text{mean}(f_{LBP_{ij}}(M > 0))$;

$F_{ij2} \leftarrow \text{std}(f_{LBP_{ij}}(M > 0))$;

$F_{ij3} \leftarrow \text{median}(f_{LBP_{ij}}(M > 0))$;

$F_{ij4} \leftarrow \text{entropy}(f_{LBP_{ij}}(M > 0))$;

$F_{ij5} \leftarrow \text{skewness}(f_{LBP_{ij}}(M > 0))$;

$F_{ij6} \leftarrow \text{kurtosis}(f_{LBP_{ij}}(M > 0))$;

$F_{ij7} \leftarrow \text{mean}(f_{VAR_{ij}}(M > 0))$;

$F_{ij8} \leftarrow \text{std}(f_{VAR_{ij}}(M > 0))$;

$F_{ij9} \leftarrow \text{median}(f_{VAR_{ij}}(M > 0))$;

$F_{ij10} \leftarrow \text{entropy}(f_{VAR_{ij}}(M > 0))$;

$F_{ij11} \leftarrow \text{skewness}(f_{VAR_{ij}}(M > 0))$;

$F_{ij12} \leftarrow \text{kurtosis}(f_{VAR_{ij}}(M > 0))$;

end

end

a wrapper method (Kohavi and John, 1997) with forward (best first)

selection. The same type of classifier is used in both the internal and external loops. The external loop divides the set into 10 non-overlapping pairs of training (90%) and test (10%) sets. For each fold of the external CV, the training set is further divided into 5 non-overlapping sets by the internal CV loop. The internal loop is done first to select the feature subset of this particular fold of the external loop. Thereafter, the external loop trains the classifier using this subset, and tests it on the remaining 10%. This is repeated for every fold. Notice that the feature set might vary with each external fold of the CV scheme. Thus doing an external or nested CV gives a measure of how well the method works for this dataset, where the method includes the feature subset selector and the choice of classifier. Figure 4.6 shows how the external CV is performed for the first fold.

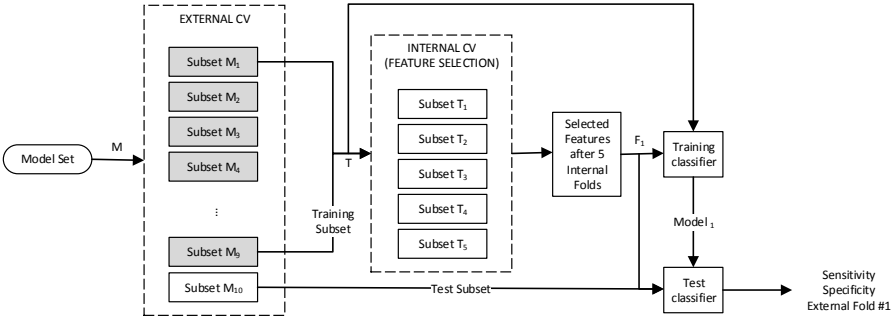


Figure 4.6: External cross validation flowchart. The process is repeated iteratively for the 10 folds of the external loop.

Finally, a final classifier is made using the whole model set for feature subset selection and thereafter the whole model set is used for training the classifier. The validation set is tested on the final classifier. The process is summarized in Figure 4.7. The normalization parameters from the model set are saved as a part of the classifier, such that the validation set is normalized using these same parameters.

Both, data preprocessing and classification, were carried out with

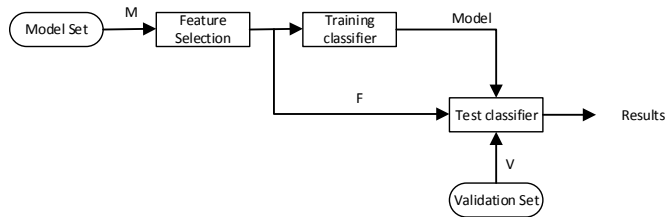


Figure 4.7: *Final validation flowchart.*

Weka (Hall et al., 2009; Witten et al., 2011).

4.4 Results

This section describes the material used and the experiments designed for method validation.

4.4.1 Material

The material used in this work are images previously diagnosed as normal (without known pathology), DR or AMD. The dataset was composed of images from 4 different databases which included some of the categories under study:

ARIA

ARIA database (Farnell et al., 2008; Zheng et al., 2012) is formed by 143 color fundus images (768×576 pixels), which are organized into three classes: age-related macular degeneration (AMD) subjects ($n=23$), healthy control-group subjects ($n=61$), and diabetic subjects ($n=59$).² Trained image analysis experts traced out the blood vessels, and also the optic disk and fovea where relevant.

²www.eyecharity.com/aria_online.html

STARE

STARE database (Hoover et al., 2000; Hoover and Goldbaum, 2003) is a full set of 402 images (700×605 pixels) where thirteen different diagnoses were considered.³ From this dataset, three subsets were generated: age-related macular degeneration (n=47), normal (n=37), and diabetic retinopathy (n=89).

E-OPHTHA

E-OPHTHA (Decencière et al., 2013) is a database of fundus images especially designed for diabetic retinopathy screening.⁴ It contains 257 images with no lesion, 47 images with exudates and 148 with microaneurysms or small hemorrhages making a total of 174 images with diabetic retinopathy.

DIAGNOS

DIAGNOS (Garnier et al., 2014) is a private database, property of DIAGNOS Inc., composed of 45 fundus images, 22 afflicted with AMD and 23 healthy.

Both E-OPHTHA and DIAGNOS have a range of different image resolutions. The four databases experience a significant variability in color, illumination, resolution, quality, etc. both within and, even more, among the databases. Some representative images of each database are shown in Figure 4.8.

All images of the resulting dataset must comply with certain quality criteria. The following causes were considered reasons for exclusion:

- Images with artefacts, for example bright and circular spots produced by some dust in the camera lens.

³www.ces.clemson.edu/~ahoover/stare/

⁴www.adcis.net/en/Download-Third-Party/E-Ophtha.html

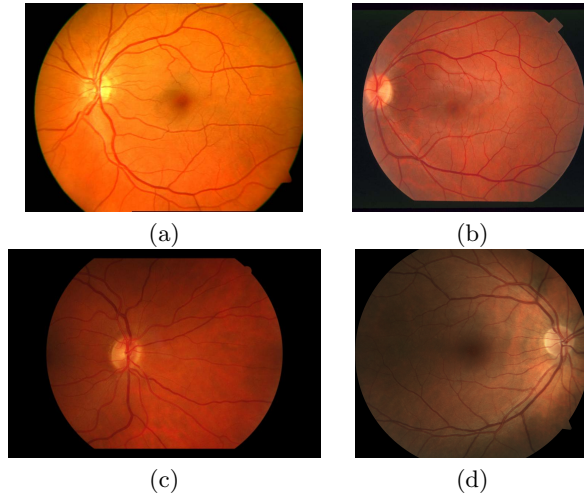


Figure 4.8: *Fundus images belonging to the different used databases. (a) ARIA, (b) STARE, (c) E-OPHTHA and (d) DIAGNOS.*

- Images affected by a relative large amount of impulsive noise (salt and pepper noise).
- Images where the vascular network is largely over-segmented by the method presented in Subsection 3.3.1.
- Images with a doubtful diagnosis.⁵
- Images with highlights around the vessels associated with young retinas.⁶
- Tessellated images due to the fact there are lesser amounts of pigment in the retinal pigment epithelium.⁶

⁵Based on second opinion from a medical doctor.

⁶This situation dramatically changes the images, thus they are not included here. However they should be regarded separately in future studies since it is highly desirable to be able to include these types of subjects.

Figure 4.9 depicts some of these cases. Most of these choices were done to determine if LBPs were able to discriminate between healthy and pathological images in a normal situation or, in other words, without the presence of distracting elements. When this hypothesis was confirmed, particular cases will be included in the analysis.

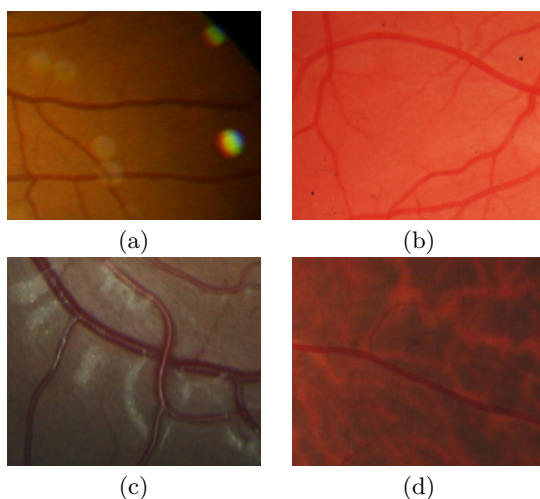


Figure 4.9: *Excluded images. (a) With artefacts, (b) Noisy, (c) With highlights and (d) Tessellated.*

After exclusion, the resulting dataset used in this work is formed by a total of 251 images. In Table 4.1, the number of images of each database after the exclusion criteria is detailed. As mentioned before, this dataset was divided into two subsets, one for training and testing by cross validation (model set) and other, completely independent, purely for testing (validation set). The model set contains 80% of the images and the validation set the remaining 20%. Table 4.2 shows the contents of each subset.

Table 4.1: *Content of each database.*

	ARIA	STARE	E-OPHTHA	DIAGNOS	<i>Total</i>
AMD	9	23	0	17	49
DR	8	35	37	0	80
NORMAL	30	13	79	0	122
<i>Total</i>	47	71	116	17	251

Table 4.2: *Content of the model and validation sets.*

	ARIA	STARE	<i>Model set</i> E-OPHTHA	DIAGNOS	<i>Total</i>
AMD	7	18	0	14	39
DR	6	28	29	0	63
NORMAL	24	10	63	0	97
<i>Total</i>	37	56	92	14	199

	ARIA	STARE	<i>Validation set</i> E-OPHTHA	DIAGNOS	<i>Total</i>
AMD	2	5	0	3	10
DR	2	7	8	0	17
NORMAL	6	3	16	0	25
<i>Total</i>	10	15	24	3	52

4.4.2 Experiments

Five experiments were conducted and validated with the procedure proposed in this chapter:

- AMD - Normal
- DR - Normal
- Pathological (including AMD and DR) - Normal
- AMD - DR
- 3 class problem (AMD - DR - Normal)

For each experiment, different classifiers were tested: Logistic Regression (Cessie and van Houwelingen, 1992), Neural Networks (Hecht-Nielsen, 1990), SVM (Chang and Lin, 2011), Naive Bayes (John and Langley, 1995), C4.5 (Quinlan, 1993), Rotation Forest (Rodriguez et al., 2006), Random Forest (Breiman, 2001), and AdaBoost (Freund and Schapire, 1996). The performance of the algorithms was evaluated based on two concepts: sensitivity or true positive rate (TPR) and specificity or true negative rate (TNR). Sensitivity and specificity measure the proportion of positive and negative cases which are correctly identified as such, respectively. Tables 4.3 and 4.4 detail the sensitivity and specificity obtained on the model and validation sets in all the experiments with the different classifiers. In the case of the three-class problem, the average sensitivity and specificity are shown. In Table 4.3, the best results of each experiment are highlighted in bold. The metrics computed after external cross validation are additive over the folds, i.e. the correctly or incorrectly classified cases are summed over the folds and divided by the total number of instances.

Table 4.3: Results of the different experiments on the model set (external CV).

	<i>AMD-Normal</i>		<i>DR-Normal</i>		<i>Path.-Normal</i>	
	TPR	TNR	TPR	TNR	TPR	TNR
Logistic Regression	0.948	0.990	0.732	0.814	0.843	0.861
Neural Networks	0.866	0.979	0.722	0.773	0.833	0.832
SVM	0.938	0.959	0.773	0.876	0.784	0.842
Naive Bayes	0.990	0.959	0.588	0.794	0.647	0.782
C4.5	0.969	0.938	0.897	0.804	0.853	0.851
Rotation Forest	0.938	0.959	0.835	0.866	0.892	0.881
Random Forest	0.959	0.959	0.876	0.866	0.843	0.861
AdaBoost	0.979	0.959	0.897	0.814	0.824	0.772

	<i>AMD-DR</i>		<i>AMD-DR-Normal</i>	
	TPR	TNR	TPR	TNR
Logistic Regression	0.873	0.937	0.835	0.918
Neural Networks	0.905	0.921	0.842	0.921
SVM	0.841	0.968	0.852	0.926
Naive Bayes	0.857	0.952	0.777	0.888
C4.5	0.889	0.937	0.849	0.924
Rotation Forest	0.921	0.937	0.883	0.942
Random Forest	0.921	0.921	0.907	0.954
AdaBoost	0.921	0.968	0.808	0.904

With regard to the obtained results after external CV (Table 4.3), the experiment with the best performance is “AMD-Normal” achieving a sensitivity and specificity greater than 0.95. The rest of the experiments let show that the lower values correspond to the detection of the DR. This is because the lesions of the DR are usually smaller than those of the AMD and, therefore, they are more difficult to detect through texture analysis. Even so, the results are promising in the three-class diagnosis.

Table 4.4: *Results of the different experiments on the validation set.*

	<i>AMD-Normal</i>		<i>DR-Normal</i>		<i>Path.-Normal</i>	
	TPR	TNR	TPR	TNR	TPR	TNR
Logistic Regression	1.000	0.920	0.412	0.960	0.815	0.920
Neural Networks	1.000	1.000	0.647	0.960	0.704	0.960
SVM	0.800	0.920	0.412	0.920	0.778	0.920
Naive Bayes	1.000	0.960	0.353	0.920	0.556	0.800
C4.5	1.000	0.960	0.824	0.880	0.741	0.880
Rotation Forest	0.900	0.920	0.706	0.840	0.815	0.880
Random Forest	1.000	0.920	0.471	0.920	0.667	0.840
AdaBoost	1.000	0.960	0.647	0.920	0.778	0.880

	<i>AMD-DR</i>		<i>AMD-DR-Normal</i>	
	TPR	TNR	TPR	TNR
Logistic Regression	0.900	0.882	0.769	0.878
Neural Networks	0.900	0.882	0.808	0.875
SVM	1.000	0.824	0.750	0.812
Naive Bayes	1.000	0.882	0.712	0.816
C4.5	1.000	0.882	0.808	0.874
Rotation Forest	1.000	0.941	0.808	0.857
Random Forest	1.000	0.824	0.808	0.892
AdaBoost	0.900	0.824	0.769	0.886

Besides the results of the external CV, an independent set of 52 images (validation set) was saved to test the proposed method. The validation results follow the same trend shown in the CV, i.e. the “AMD-Normal” experiment is the one which achieves the best performance.

The proposed method was also compared with other state-of-the-art algorithms which are based on fundus image texture analysis ([Garnier et al., 2014](#); [Mookiah et al., 2013](#)). As mentioned in the introduction, Garnier *et al.* use the LBP histogram calculated after a wavelet decomposition as input of a linear classifier validated by means of the leave-one-out method. Mookiah *et al.* combine the LBP information with other features (the area of hard exudates, the area of blood vessels, bifurcation points, etc.) in a probabilistic neural network. To the best

of the author's knowledge, there is no other system that analyses the texture of the retina background and detects AMD and DR at the same time, therefore, it was only possible to compare the results from two-class diagnosis, see Tables 4.5 and 4.6. The results of the proposed method shown in these tables are those achieved on the model set because the other compared works do not test their algorithms on an independent dataset. Note that the obtained results are not directly comparable because the dataset used in each case is different.

Table 4.5: *AMD-Normal comparison on different databases.*

	TPR	TNR	Features	Classifier	Validation	#images
Garnier <i>et al.</i>	0.913	0.955	Wavelet transform and LBP	LDA	Leave-one-out	45
Proposed method	0.979	0.959	LBP	AdaBoost	10-fold external CV	136

Table 4.6: *DR-Normal comparison on different databases.*

	TPR	TNR	Features	Classifier	Validation	#images
Mookiah <i>et al.</i>	0.963	0.961	Blood vessel area, exudates area, bifurcation points, texture and entropy	PNN	3-fold CV	156
Proposed method	0.876	0.866	LBP	Random Forest	10-fold external CV	160

4.5 Conclusions

In this chapter, a new approach for AMD and DR diagnosis has been presented. It is based on analyzing texture discrimination capabilities in fundus images to discriminate healthy patients from images with AMD or DR. The performance of LBP with different classifiers was tested.

Many more AMD or DR detection techniques exist in the literature but most of them focus on lesion segmentation instead of a direct study of the retina background. This fact makes the accuracy of the classification stage dependent on the accuracy of the lesion segmentation. Lesion segmentation involves a series of uncertainties and a non-accurate segmentation may provoke important errors in the classification. The main advantage of the proposed procedure is that it gets a good performance without having to search different types of lesions. The only segmentation needed in the presented approach is to mask the significant structures (vessels and optic disk) but their accuracy has little influence on the final result.

The obtained results, a sensitivity and specificity greater than 0.96 for AMD detection, greater than 0.87 for DR detection and greater than 0.91 for three-class diagnosis, demonstrate that LBP is a good texture descriptor for fundus images and can be used in a diagnosis aid system for retinal disease screening.

As for future work lines, more images could be added to the database and the analysis could be repeated. Moreover, some work should be carried out to develop strategies that enable the analysis of the type of images that were excluded from the initial database, such as tessellated fundus, images with highlights or typical artefacts.

Chapter 5

Conclusions

As was indicated in Chapter 1, the main objective of this thesis was to develop and implement algorithms for retina characterization from the analysis of fundus images. In particular, the retina was characterized through advanced image processing techniques in two different ways. First, the main retinal structures, blood vessels and optic disk, were detected and subsequently analyzed to obtain some of their significant features. This information was used in a computer-aided diagnosis software for hypertensive risk determination. This was addressed in Chapter 3. Secondly, the texture of the retina background was studied in order to classify an image as belonging to a healthy or pathological subject, as was seen in Chapter 4. Diabetic retinopathy and age-related macular degeneration were detected by this procedure. Its main advantage is to avoid the need for segmentation of each type of retinal lesion.

On the one hand, Chapter 3 demonstrated that the performance of mathematical morphology and stochastic watershed, a variant of the watershed transformation, allows us to apply these methods for clinical purposes as the segmentation of the retinal structures. On the

other hand, Chapter 4 stood out that local binary pattern operator is a good texture descriptor for fundus images and can be used for retinal disease screening, above all for the diagnosis of age-related macular degeneration.

The methods presented in this thesis will be incorporated in an only automatic screening system to discriminate between a normal and pathological retina taking into account different pathologies at the same time. This system will have to be more robust than the existing solutions and to have enough specificity and sensitivity to be applied in clinical practice. Nowadays, there are some initiatives focused on glaucoma or diabetic retinopathy detection but they have very low specificity and, therefore, low screening effectiveness. The development of this screening system is being funded at the present time by the Spanish government under the project ACRIMA (TIN2013-46751-R). It is expected to have a first version of the system at the end of 2016. In this project, new prognostic markers that are currently used in clinical diagnosis, such as blood pressure, age or family history, will be combined with those provided by the fundus image processing with the aim of improving the robustness and increasing the sensitivity and specificity of the automatic screening system. From a perspective of a screening tool, all developments presented in this thesis should be validated by comparing the results achieved by the tool with the diagnosis made by an expert.

In future works, three main research lines will be carried out. First, to segment other retinal structures such as the macula or the optic cup. The location of the macula is important for the diagnosis of age-related macular degeneration, as its name suggests. The area of the optic cup is directly related to glaucoma prediction, as was seen in Chapter 2. Secondly, to perform an in-depth study of the retina background. More texture descriptors will be evaluated and their results will be compared with those presented in Chapter 4. Finally, to consider the heterogeneity of the clinical databases. The parameters used in the developed methods which are related to the image size will be generalized. The diverse

types of normality that exist in fundus images, as the tessellated images, will have to be taken into account to not reduce the specificity of the classification, which is increasingly important.

Merits

Journal papers

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