

Automated Angiogenesis Quantification through advanced Image Processing Techniques

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Abstract—Angiogenesis, the formation of blood vessels in tumors, is an interactive process between tumor, endothelial and stromal cells in order to create a network for oxygen and nutrients supply, necessary for tumor growth. According to this, angiogenic activity is considered a suitable method for both tumor growth or inhibition detection. The angiogenic potential is usually estimated by counting the number of blood vessels in particular sections. One of the most popular assay tissues to study the angiogenesis phenomenon is the developing chick embryo and its chorioallantoic membrane (CAM), which is a highly vascular structure lining the inner surface of the egg shell. The aim of this study was to develop and validate an automated image analysis method that would give an unbiased quantification of the micro-vessel density and growth in angiogenic CAM images. The presented method has been validated by comparing automated results to manual counts over a series of digital chick embryo photos. The results indicate the high accuracy of the tool, which has been thus extensively used for tumor growth detection at different stages of embryonic development.

I. INTRODUCTION

Angiogenesis is a tightly regulated process that is essential for embryonic development that has also been implicated in physiological, as well as pathological phenomena in fully developed organisms [1]. Soluble growth factors, adhesion molecules, proteases, as well as the extracellular matrix have been shown to interact spatially and temporally to co-ordinate the formation of new blood vessels. Stimulation of new blood vessel growth has been proposed as a therapeutic approach to treat conditions that include ischemic heart disease, neurodegenerative disorders and hair loss. On the other hand, an excessive and deregulated angiogenic response is thought to contribute to cancer, and other diseases [2]. More precisely, the angiogenetic formation of blood vessels in tumors, is an interactive process between tumor, endothelial and stromal cells in order to create a network for oxygen and nutrients supply, necessary for tumor growth. One of the most popular assay tissues to study the angiogenesis phenomenon

is the developing chick embryo and its chorioallantoic membrane (CAM). CAM is a highly vascular structure lining the inner surface of the egg shell. The study of the formation of its vessels, at different stages of development, is efficiently indicative for the phenomenon ([7]-[12]). The aim of this study was to test and develop an automated image analysis method that would give an unbiased quantification of the micro-vessel density. The developed tool processes digital images of in vivo chick embryo and its CAM. Using advanced image processing techniques, the proposed tool exploits vessel characteristics, such as vessel length, density, vessel branching points and textural information to assess the level of angiogenesis.. The presented method has been validated by comparing automated results to manual expert assessment over a series of digital chick embryo images. The results indicate the high accuracy and reproducibility of the tool, which has been thus extensively used for tumor growth detection at different stages of embryonic development. The rest of the paper is organized as follows; Section 2 presents related work regarding CAM development and automated angiogenesis quantification. Section 3 describes the CAM angiogenesis procedure and Section 4 presents the proposed quantification methodology. Evaluation results are presented while, Section 6 concludes the paper.

II. RELATED WORK

CAM is one of the most widely used models to study angiogenesis. Attempts to quantify angiogenesis using biochemical measurements have focused on [14C] - proline and hemoglobin determination. Type IV collagen, is abundantly expressed in the basement membrane, a structure that develops below and supports the vascular endothelium. Like all types of collagen, type IV collagen has high proline content. This method is used by a few investigators with some success [3]. Other laboratories use hemoglobin content as a marker for angiogenesis [4]. Most investigators prefer to quantify the angiogenic response in the CAM by manual scoring or by morphometric analysis of CAM pictures using appropriate software, such as Scion Image [5], or Image-Pro Plus [6]. The latter tools provide image processing techniques, which allow measurements of vascular length after being applied manually on the examined angiographic images. Additional work in literature ([9]-[12]) presents computer-aided methods for angiogenesis quantification

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which involve manual processing of the images for calculating vessel density, vessel length and branching points. To our knowledge there is no automated software for angiogenesis quantification currently available. In this context, the presented method provides an automated angiogenesis assessment based on vessel length, branching points and texture quantification. Additional features allow the calculation of vessel thickness and the automated process of numerous experimental images.

I. EXPERIMENTAL SETUP AND IMAGE ACQUISITION

This section describes how CAM and corresponding digital images are acquired for the experiment. Fertilised White Leghorn chicken eggs were placed in an incubator and kept under constant humidity at 37°C. On day 4, a square window was opened in the shell and then sealed with adhesive tape. In experiments designed to monitor the development of vascular networks at different stages of development, tissues were collected on day 9, 10, 11 or 12. In a separate series of experiments we evaluated the effects of dexamethasone, a known angiogenesis inhibitor. For these experiments, an O-ring (1 cm²) was placed on the surface of the CAM on day 9 and dexamethasone (8nmol/egg) was placed inside this restricted area. After 48 hr, CAMs were fixed in Carson's solution (saline-buffered formalin) and color images of the collected tissues were obtained through a Canon Powershot A620 digital camera (7.1 MPixels) using a Carl Zeiss W10X/20 stereoscope (6.3–25mm object field, up to 32x zoom). Obtained images were processed by the proposed image analysis tool presented in the next section that automates the assessment of the angiogenic procedure.

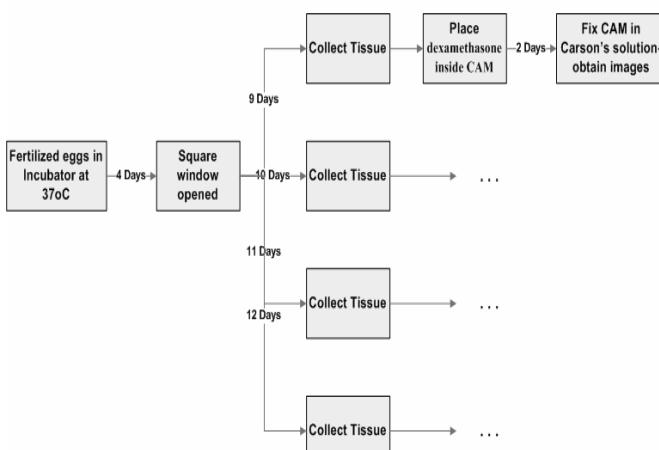


Figure 1. Experimental setup and image acquisition procedure

An illustration of the aforementioned experimental procedure is provided by Figure 1.

II. AUTOMATED ASSESSMENT OF ANGIOGENESIS IN STEREOSCOPIC IMAGES

The automated processing and angiogenesis quantification of the stereoscopic images is performed in three basic phases (see Figure 2); during phase A the angiogenic image is captured and converted to grayscale. Phase B performs adaptive thresholding and noise removal. During adaptive thresholding, a user predefined size of pixel neighborhood is processed and the mean value M is calculated. Then for every pixel value p , if $p < M - C$, where C is a user defined constant, the p is set to the background value. Otherwise, the foreground value is assumed. Noise removal follows using a Median Filter adapted from [13]. Additional noise removal techniques found in literature include conservative smoothing [14], Gaussian filtering [15], Mean filtering [16] and others [17]. Median filtering has been selected due its simplicity and low complexity, resulting in faster image processing, and efficient performance on the applied sample images. The image output of phase B is illustrated in Figure 3(b). Depending on the image initial quality (lighting conditions, vessel density, tissue quality), different values of the neighborhood size and the constant C can be selected for optimizing the output image quality of phase B. During our experiments the value of C that optimizes the thresholding procedure was $C = 15$ with a neighborhood size of 5x5 pixels. Comparing Figure 3(a) and Figure 3(b), it can be obvious that issues like bad lighting effects and tissue margins on the image corners are efficiently addressed by the adaptive thresholding procedure. Phase C skeletonizes the image and produces the final image output (Figure 3(c)) that contains vessels with thickness of one pixel. Finally, the developed tool calculates automatically the total vessel length, the vessel branching points, the vessel density over the whole image area and the vessel texture, assessing this way the angiogenesis phenomenon.

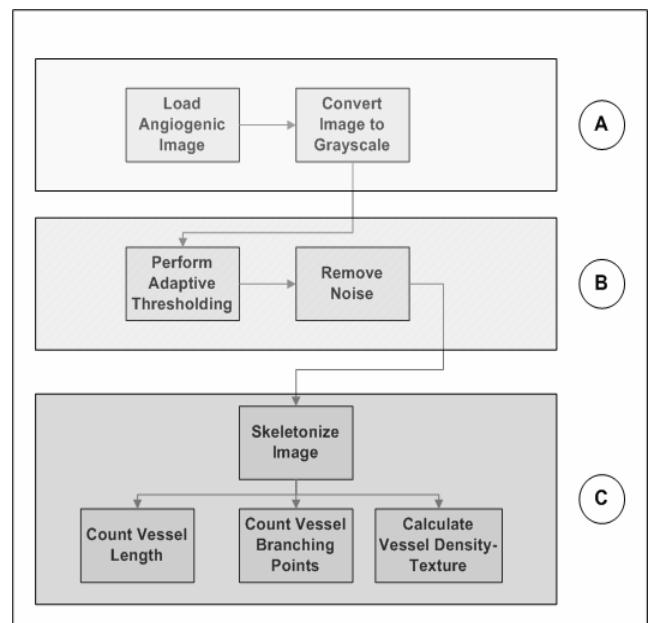
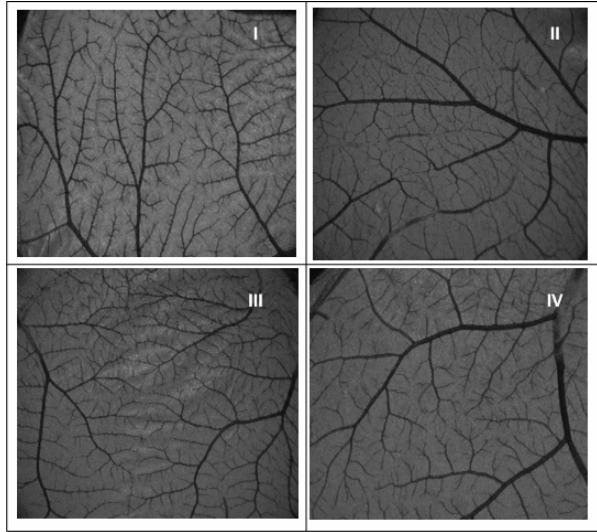
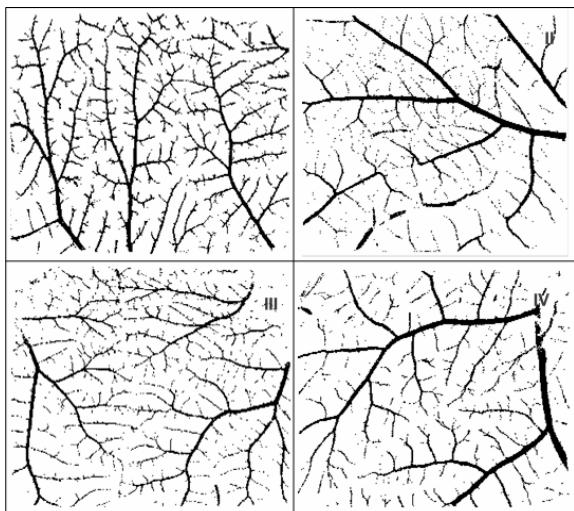


Figure 2. Block Diagram of the automated angiogenesis assessment

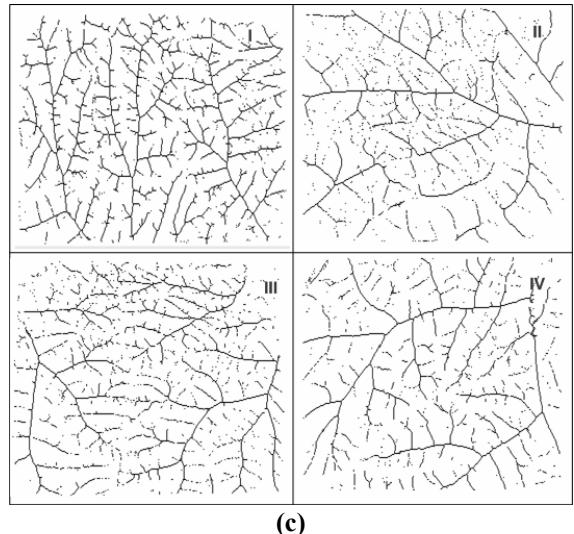
The aforementioned quantitative results are saved in XML files that can be directive saved into a database for processing. In addition, the tool offers automatic process of numerous image files defined by user, and basic image manipulation functions, like zooming, Region of Interest (ROI) selection and manual vessel thickness calculation.



(a)



(b)



(c)

Figure 3. (a) Original angiogenic CAM images after greyscale conversion, (b) images produced after adaptive thresholding and noise removal, (c) skeletonized images

The corresponding results (i.e. vessel length, vessel density, branching points and image texture in terms of contrast, correlation and entropy) for the images included in Figure 3 are presented in Table I.

TABLE I
EXPERIMENTAL ANGIOGENESIS QUANTIFICATION RESULTS

| Vessel Quantification Metric | Value | | | |
|-------------------------------|-------|-------|-------|-------|
| Image No. | I | II | III | IV |
| Vessel Length (pixels) | 6918 | 8959 | 8379 | 5877 |
| Vessel density (%) | 8.6 | 11.1 | 10.4 | 7.3 |
| Vessel Branching points | 439 | 485 | 557 | 303 |
| <i>Image Texture Features</i> | | | | |
| Contrast | 0.122 | 0.165 | 0.140 | 0.103 |
| Correlation | 4.028 | 3.025 | 3.542 | 4.565 |
| Entropy | 0.780 | 0.930 | 0.833 | 0.736 |

III. EVALUATION RESULTS AND DISCUSSION

The presented automated angiogenesis quantification method has been validated against manual counts of vessel branching points over a series (25) of angiogenic CAM images acquired with the procedure described in Section 3. The corresponding results are illustrated in Figure 4, which presents the normalized number of vessel branching points calculated or manually counted in each image divided by the mean value of branches in the whole image dataset.

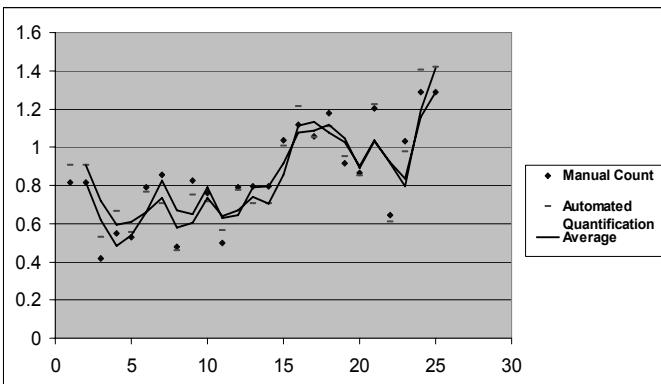


Figure 4. Automated Quantification validation results against manual counts over 25 CAM angiogenic images. Y axis represents the normalized number of vessel branches calculated or manually counted for each image, divided by the mean value of branching points in all images.

The Mean Average Error for the automated quantification of the experimental images against the manual count of branching points was 5% with an average standard deviation of 7%. The results indicate the high accuracy of the presented method in automated quantification of the vessel branching points. The latter conclusion can also avouch the accuracy of the additional calculated features (i.e. the vessel length, density and texture). The tool has been thus extensively used by biologists for quantification of the angiogenesis phenomenon and tumor growth detection and inhibition.

IV. CONCLUSION

Automated quantification of angiogenesis levels is quite important in the objective assessment of drugs, chemical substances or anti-angiogenic procedures in general. The current methodology used by several medical labs is based on either on the visual count or in the use of commercially available image analysis SW (i.e. Scion Image, Image-Pro Plus, etc.). This methodology introduces large variation, caused by inter- or intra-operator sensitivity regarding the visual counts, or different setup of parameters and various processing steps regarding the existing image analysis SW. Quite important is also the standardization of image acquisition procedure which ensures the reproducibility of the captured images.

In this study, we aim at the implementation of a fully automated, objective, computer-based image analysis tool for the determination of angiogenesis level in digital stereoscopic images. At this stage we demonstrated that the implemented tool can calculate correctly several features, related to quantification of angiogenesis, such as vessel length, density and branching points. The produced results were highly correlated with the average human visual counts, conducted by experts from the Institute of Biological Research and Biotechnology in National Hellenic Research Foundation and in the Lab of Molecular Pharmacology, School of Pharmacy, University of Patras in Greece. Our future work involves the introduction of advanced classifiers in the proposed tool (Statistical, Neural Networks, Support

Vector Machines, etc) and the exploitation of various image textural features for achieving further quantification and automation in the assessment of angiogenesis level.

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