ANALYSIS OF DUAL-ENERGY X-RAY ABSORPTIOMETRY IMAGES USING COMPUTER VISION METHODS

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ABSTRACT
PURPOSE: Dual-energy x-ray absorptiometry (DXA) is the “golden standard” for diagnosing osteoporosis. Its analyzing algorithm (software) makes it possible to distinguish the bone from the soft tissue. Until now there are only attempts to process and acquire images using automatic segmentation with convolutional neural networks (CNN). Machine reconstruction and precise specific models of anatomic structures from medical images could be accomplished using computer vision. The objective of the current work is to introduce the potential of the two computer methods and their application in the diagnostic DXA analysis.

METHODS: DXA generates a report in the DICOM format which includes patient data (age, gender, height, weight, bone mineral density, T-score and Z-score) and an image of the scanned spine as well as the region of interest (ROI). The CNN methods are based mainly on intermediate analysis. The learning of the segmentation of CNN by generating segmentation labels using simple heuristic is done using computer vision. The functions of the loss and the architecture of the CNN is then determined. In that manner the right analysis of the existing medical image is made possible.

RESULTS: The computer library OpenCV is the way to realize a model for the assessment of a DXA analysis. The library is available for Python programming language. The library has functions for the extraction of colour objects, image smoothing, Canny’s edge detector, Hough transform and methods for work with contours.

CONCLUSIONS: The detection and extraction of images is fundamental for the analysis of DXA which is a step forward in the precision of the in-vivo diagnostic of the bone.

Key words: Dual-energy x-ray absorptiometry (DXA), convolutional neuronal networks, computer vision, diagnostic analysis

PURPOSE
The analysis of medical images requires a combination of interdisciplinary fields, including electrical, computer and biomedical engineering, as well as other fields (1). Despite the independent development of medical imaging and computer vision, the merging of these two sciences holds promising results. Scientific papers focused on the analysis of medical images with computer vision were presented at the European Conference on Computer Vision (ECCV). In 2006, a seminar on “Approaches of computer vision to the analysis of medical images” was held in Graz, Austria (2). The application of computer vision techniques for the analysis of medical images is used for both clinical evaluation and operations. Automated reconstruction of precise specific models of anatomical structures from medical images is becoming essential for surgical procedures, disease research, and for clinical evaluation of the effectiveness of therapy (3, 4). For example, in minimally invasive procedures, neurosurgeons often use image-related navigation aids to locate the field of operation (5). Therefore, the extraction of anatomical models from images by computer vision is a key element of emerging surgical techniques (6).

The only DXA image analysis has been done with a Convolutional Neural Network (CNN) (7). This methodology is based on intermediate analyzes. They can improve classification efficiency in the context of specific medical
applications, as they are trained by databases that are usually available in the medical image. Interim analyzes incorporate prior knowledge of how pathology is imaged on DXA and provides important guidance for the network. The goal is to automatically segment the marked body parts of the DXA image with a Convolutional Neural Network (CNN). The segmenting CNN receives the DXA image at the input and produces six different channels with the same dimension as the input, where each channel corresponds to the six marked parts. Because DXA scanning of the whole body is homogeneous, segmentation labels for some parts of the body can be produced with a series of simple heuristics. These labels, while not perfect, are good enough to train the segmenting CNN, and in many cases, the trained CNN produces visually better segmentation. To date, no specific analyzer of DXA test results by computer vision has been developed.

The DXA study generates a report, which is in DICOM format and includes patient data (age, sex, height, weight, bone mineral density, T-score and Z-score) and an image of the scanned spine, as well as location of the field of interest.

The objective of the current work is to introduce the potential of the two computer methods and their application in the diagnostic DXA analysis.

METHODS

The following methods of computer vision were used to perform the set tasks: extraction of colored objects from an image, Gaussian blurring, detection of boundaries in an image (Canny's algorithm), Hough transformation and contour detection. The file format of the examined data was the DICOM standard.

Each DICOM report from DXA scan includes:
- patient data (age, gender, height, weight, bone mineral density (BMD), T-score and Z-score)
- image of the scanned spine as well as the region of interest (ROI).

The program implementation of the methods was performed in Python programming language through the libraries PyDICOM, OpenCV, NumPy and PIL. The Tkinter module was used to create the graphical interface of the application.

Once detected, each of the features can be extracted from the DXA scan image. This extraction often requires the use of a large amount of resources and the result is known as a descriptor or attribute vector.

Feature detection involves methods for calculating image information and making decisions at each point in the image for the presence of a feature of a particular type. Feature extraction is a dimensional reduction process that starts with the original set of output variables, from which less is constructed for accessible processing and sufficiency to accurately and completely describe the original data set. Sign extraction is used in machine learning, image recognition and image processing. It starts with the initial set of data from which we obtain secondary derivative values (signs), which are considered to be informative and unambiguous. This achieves a consistent process of machine learning and generalization of the steps and
leads to a better human interpretation of the data. One of the most important areas of application is image processing, in which algorithms are used to detect and isolate a variety of desired areas or shapes (characteristics) of a digital image. It is widely used in Optical Character Recognition (OCR) (15).

RESULTS
The OpenCV computer library is used for the software implementation of the DXA analysis. It is available in Python under the name "cv2" and is included in the program with import cv2. The library uses functions for program implementation of the method for extracting colored objects, the method for image blurring, the Canny algorithm, the Hough transformation and the methods for working with contours. The extraction of colored objects is required in the evaluation for the correct location of the region of interest. Initially, the color space of the image is changed with cvtColor (input_image, flag). Then, by setting the color boundary values of the field of interest, the inRange (input_image (HSV), lower_bound, upper_bound) function is used by OpenCV, extracting only its pixels. The bitwise_and function (input_image, output_image, mask = mask) masks the field of interest from the original image and extracts it. The program implementation of the process is part of the program method of the analyze_roi () model.

Gaussian blur is applied in the computer model to evaluate DXA analysis in order to preprocess the image and to reduce noise before applying other methods. It was used before Canny's algorithm and Hough's transformation. In Python, Gaussian blur is implemented with the function from the OpenCV GaussianBlur library (input_image, (matrix size)) and is performed in the program method of the model analyze_angle () and analyze_roi (). The result is an image with a lower noise level and clearer borders. This improves the performance of Canny's algorithm, which detects boundaries more efficiently, which in turn is needed for better Hough transformation results. The Canny algorithm is the main method for detection of boundaries in the image used in the DXA analysis evaluation model. The application of the algorithm is performed in the analysis to calculate the distortion angle before the Hough transformation and before the detection of the contours of the field of interest.

Hough transformation, as one of the most effective methods for detecting lines in an image, is used in the DXA analysis evaluation model. The detection of straight lines is performed when calculating the angle of distortion. The aim is to find the lines corresponding to the vertebrae and to calculate the angle of curvature. The program implementation of the method in Python is performed through the function from the OpenCV HoughLines library (input_image, ρ, θ, threshold) and is part of the program method of the analyze_angle () model.

DISCUSSION
Using the methods of computer vision algorithm was developed to analyze DXA scan images. Based on the developed application for analysis of DXA images, important information can be derived about the angle of curvature in the lumbar region in people who have performed a DXA scan. The application may be useful for conducting large-scale studies to determine the incidence of spinal curvature from DXA images, as this type of study involves less radiation than a standard X-ray. Additionally, the application helps to correctly interpret the DXA scans, as it provides information about the angle of lumbar curvature. To date, no such DXA image analyser based on computer vision has been developed, and this thesis lays the foundations for a significant discovery. This application can be used in the future in medical practices in order to improve the diagnosis of osteoporosis.

CONCLUSIONS
The detection and extraction of images is fundamental for the analysis of DXA which is a step forward in the precision of the in-vivo diagnostic of the bone.

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