Three-dimensional periodontal investigations using a prototype handheld ultrasound scanner with spatial positioning reading sensor

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Abstract

Aim: To demonstrate the feasibility of the 3D ultrasound periodontal tissue reconstruction of the lateral area of a porcine mandible using standard 2D ultrasound equipment and spatial positioning reading sensors. Material and method: Periodontal 3D reconstructions were performed using a free-hand prototype based on a 2D US scanner and a spatial positioning reading sensor. For automated data processing, deep learning algorithms were implemented and trained using semi-automatically segmented images by highly specialized imaging professionals. Results: US probe movement analysis showed that non-parallel 2D frames were acquired during the scanning procedure. Comparing 3 different 3D periodontal reconstructions of the same porcine mandible, the accuracy ranged between 0.179 mm and 0.235 mm. Conclusion: The present study demonstrated the diagnostic potential of 3D reconstruction using a free-hand 2D US scanner with spatial positioning readings. The use of automated data processing with deep learning algorithms makes the process practical in the clinical environment for assessment of periodontal soft tissues.

Keywords: periodontal ultrasonography; 3D imaging; artificial intelligence; ultrasound imaging; periodontal diagnosis

Introduction

Periodontal probing is the accepted gold-standard approach for the assessment of the periodontal soft-tissue [1]. Noninvasive techniques to image gingival soft tissues are not currently available in dentistry. Bi-dimensional ultrasound (2D US) imaging has already shown great potential for evaluating periodontal soft-tissue structures [2-5]. Lin et al demonstrated that periodontal probing depths could be measured with photoacoustic imaging [6]. The imaging modality was more precise compared with periodontal probing, offered higher resolution images and covered all areas of the tooth. The gingival thickness could also be easily measured using ultrasound [7].

However, 2D periodontal US is not commonly used because clinicians have to imagine the three-dimensional (3D) view of the anatomic structures within the tissue volume from the planar US images [8]. Mechanical 3D probes are relatively compact and are convenient to operate, though they are larger than conventional linear probes. Their data acquisition and reconstruction times are short, which allows the viewing of high-quality 3D images in real-time. Volumetric US systems can fully capture a 3D region of interest, but current 3D systems require specialized transducers which are prohibitively expensive for many clinical departments and do not register image orientation with respect to the patient. Clini-
cians are required to hold the mechanical 3D probes statically while acquiring images, which can lead to latent errors for data acquisition [9].

Developing smart computer algorithms to reconstruct 3D data volume using 2D frames could overcome problems related to 2D US imaging such as low reproducibility and operator dependency. Moore et al used a photoacoustic US transducer connected to a stepper motor for axial scanning and a sliding positioner for lateral control to obtain 3D images [10].

For 3D US imaging, besides the positioning and orientation errors that may corrupt the reconstruction process, another source of operator’s dependency is image interpretation. Compared with other medical images (X-ray, CT, MRI), the US images are characterized by higher noise, image contrast dependent on tissue density and low reproducibility. The interpretation is time consuming and is strongly dependent on clinician’s experience. To overcome these challenges, various semi-automatic and automatic techniques to analyze the US images have been developed. The traditional approach was to use hand-crafted image processing techniques, such as edge detection, contrast enhancement, or line fitting [11,12].

The development of deep learning models for computer vision has changed the paradigm for automatic US image processing and analysis. Deep learning is a powerful tool for extracting nonlinear features from data. This is particularly promising in US imaging, where predictable acoustic patterns are typically neither obvious nor easily hand-engineered [13].

To obtain the spatial orientation of the acquired 2D US periodontal images with respect to the patient and to generate 3D US model reconstruction of a larger area such as a maxillary hemiarch, our team developed a handheld ultrasound scanner prototype based on a standard 2D US machine and a spatial positioning reading sensor. In this paper, we report the preliminary results of periodontal 3D reconstruction using the prototype. For automated data processing, deep learning algorithms were implemented and trained using semi-automatically segmented images by highly specialized imaging professionals. The main objective of this study was to demonstrate the feasibility of the 3D US periodontal tissue reconstruction of the lateral area of a porcine mandible. The secondary objective was the evaluation of the accuracy and reproducibility of the method.

**Material and method**

A 3D high frequency US imaging prototype based on a standard medical US scanner (Vinno 6, Suzhou, China) with a high frequency (10-23 MHz) and small aperture (12.8 mm) linear transducer (X10-23L, Vinno, Suzhou, China) and a positioning reading sensor embedded in an articulated measurement arm (Evo 7, RPS Metrology, Sona, Italy) was developed by Chifor Research SRL team and used for the study. The probe was attached to the sensor and the scanning was performed at 23 MHz. The positioning reading sensor tracked the trajectory of the transducer, providing coordinates and orientation during scanning as the output. The scanner had a free-hand architecture with 6 degrees of freedom. The transducer and the sensor were position calibrated and time synchronized at the beginning of the first scanning procedure.

To test the accuracy and reproducibility of the prototype, three consecutive US scans (Scan 1, Scan 2 and Scan 3) were performed on the lingual side of 4 lateral teeth (two molars and two premolars) of a porcine mandible immersed in water (fig 1). The US examinations of the pig mandible were performed in January 2020 by the same operator. The mandible was harvested from a pig raised and slaughtered for alimentary purposes in farms organized according to European Union legislation and was not from experimental animals. The scans started between the third and second molars, moved in mesial direction to the second premolar and then moved backwards, in distal direction to the starting point. The US scanner had a frame rate of 30 frames or images per second; 500 consecutive frames from each scan were used for 3D reconstruction corresponding to the lateral area of the mandible. A total number of 1500 2D periodontal US images were taken into consideration for the prototype accuracy evaluation. For each 2D frame, the spatial and orientation information was utilized to organize the frame in the 3D space.

**Semi-automatic segmentation**

Prior to the 3D reconstruction, the 2D images were segmented for three types of tissues, i.e., tooth, bone and gingiva. The 2D images of Scan 1 were annotated and segmented by the research team at the University of Alberta using the in-house developed labeling software. Figure 2 shows the workflow of the segmentation process to segment the alveolar bones automatically. An oral US imaging specialist or reader initiated a graph-cut based segmentation process by roughly defining a region inside
the desired structures in selected frames to serve as seed pixels. This process started off in a frame and was skipped for the next few frames if the consecutive frames had similar structures. It was then followed by an automatic contour interpolation and refinement process, which used an active contour model to incorporate the intensity and shape prior information from the previous step. The reader checked the interpolation result and revised the labeling manually with the built-in “brush” and “eraser” tools to ensure accurate segmentation for every frame.

To extract the Region of Interest (ROI), representing the marginal periodontal tissue area (cortical bone, tooth crown and gingival tissue) and to exclude the other part of the image, the integral projection along the horizontal axis of the images upon the depth axis was performed and modified by multiplying the integral curve with a one-side Tukey taper, which did not alter the early portion of the curve but suppressed the tail of the integration. The result-ant was a bimodal signal which represented the ROI and the structure-free region, respectively. The Otsu’s thresholding method [14] was then used to find the optimum position separating these two peaks to extract the ROI.

The simple linear iterative clustering (SLIC) technique [15] was applied to the ROI image to group pixels with the same characteristics into superpixels for segmentation. Following Boykov et al [16], a graph-cut method was implemented to segment tissues using the seed pixels selected by the reader. The distance penalties were added to the regional cost comparing to the original graph-cut method proposed by Boykov et al [16], which was based on the observation that intensity distribution and gradient alone could not sufficiently generate accurate segmentation in our case.

The graph-cut segmentation was performed for a few frames and then the intermediate similar frames were automatically interpolated. An active contour model [17] was used to incorporate the shape and intensity prior information from the previous semi-automatic segmentation step to perform the interpolation. The in-house developed GUI software for semi-automatic segmentation was implemented in the Python programming language, based mainly on libraries such as OpenCV, Numpy, and Tkinter. The pyMaxflow library [18] was also employed to optimize the Max-flow/Min-cut algorithm for the graph-cut [17].

Automatic segmentation

Images from Scan 2 and Scan 3 were segmented by the research team at the Romanian Institute of Science and Technology using Neural Networks Algorithms and a deep learning model for object detection and image segmentation developed by extending the Mask R-CNN model [19,20]. The model was trained on an Intel® Core™ i7-9700K CPU @ 3.60GHz × 8 cores, 32GB RAM computer equipped with GeForce RTX NVidia GPU with 8GB RAM. The 500 annotated US images from Scan 1 were used for training. The annotation consists of separate images that contain the masks for 3 types of tissue: tooth, bone, and gingiva. This model was preloaded with the COCO weights [21] and was trained using the images from Scan 1. The training dataset was created by randomly selecting 420 out of 500 images from Scan 1. The other 80 images were used for model validation. The training was done on one GPU (CUDA) processing simultaneously 4 images. All Mask R-CNN weights were retrained to detect 3 new classes by running 5 epochs and 150 steps per epoch. As a result, the trained model was able to automatically detect and segment the three types of tissues.

3D reconstruction and statistical analysis

Data acquisition and 3D reconstruction were performed by the team at Chifor Research SRL using the 3D US scanner prototype and the developed software. After the US data was acquired, each frame was paired or matched with the sensor’s readings. The spatial coordinates and orientation of each frame were determined. All the frames were then re-organized in the 3D space, thus reconstructing the 3D data volume. The Cloud Compare Open-Source Software (CCOSS) Version 2.6.3 was used to compare the 3D reconstructed frames after each scan. Standard deviation (std) and mean distances were calculated between the point clouds.

Results

Table I summarizes the pertinent information about the three scans and the reconstructed data volume.

Using Scan 1 readings as example, the overall translation movement (fig 3a) shows that the US probe moved...
in a well-defined plane in one direction from distal to mesial from the 1st frame to the 295th frame and then in the opposite direction from mesial to distal from 296th to 500th frame. The relative movement of the probe between frames (fig 3b) shows the short distance pattern where the spikes represent changes in the probe movement direction between two adjacent frames. The figure also shows the long-distance movement pattern from molar 3 towards premolar 2 with two major changes in direction along the overall trajectory of the probe. While moving along the posterior-anterior direction away from the starting point, the majority of the motion are in the positive range from 1st to 295th frame. During the anterior-posterior motion, the probe moved in the opposite direction and therefore, from 296th until 500th most values are negative. Figure 3c shows the angular variations of the probe between neighboring frames, where the spikes represent changes in plane angulation movement, showing that non-parallel 2D frames were acquired during the scanning procedure.

Five hundred US images from Scan 1, which included the tooth, alveolar bone and gingiva, were segmented semi-automatically. The average speed for the semi-automatic segmentation algorithm was around 45 seconds per image. These images were scaled and converted to a specific binary format compatible with a Mask R-CNN deep learning model. Four hundred and twenty images were used as a training set and the rest of the images as a validation set. The accuracy of the training and validation sets was evaluated by the mean accuracy precision mAP [20]. For 5 epochs with 150 steps and batch size of 4, the training time was 2 hours and 40 minutes, and mAP was over 90%. At the pixel level, the precision between the annotated and detected masks was calculated as IoU (Intersection over Union). The average IoU was 85%. Scan 2 and Scan 3 were segmented using the trained model from Scan 1. All three types of tissues were identified in all images from Scan 2 and Scan 3. The precision of the automated segmentation was evaluated globally by comparing the results with those obtained in Scan 1. An example of the segmentation results is shown in Figure 4. The automatic segmentation took 0.45 second per image.

The 3D reconstruction of the teeth for the three scans is shown in Figure 5. Grey scale intensity values of the voxels from the point clouds were converted to RGB
(red, green, blue) values for better visualization. The color conversion and visualization of the point clouds were performed using CCOSS. Six arbitrary points (R0, R1, …, R5) and six corresponding points (A0, A1, …, A5) were selected by the operator in the reconstructed Scan 1 and Scan 2 areas, respectively. Using these points, CCOSS aligned the point clouds for statistical analysis to determine the mean distance and the standard deviation between them (fig 6).

Figure 7 shows the aligned 3D reconstruction of the teeth from Scan 2 and Scan 1. The 3D reconstruction was colored according to the distances between the 2 point clouds in figure 6. The mean distance and standard deviation between the points from the two scans (at tooth level) are 0.179 and 0.187 mm, respectively. Similarly, figure 8 shows the aligned reconstruction of the teeth from Scan 3 and Scan 1. Scan 3 included an extra tooth distally, which was not covered in Scan 1. The color an-

**Fig 4.** Image segmentation: (a) the original US image, (b) the semi-automatic segmentation of the image, (c) the automatic segmentation of the image, and (d) the difference between semi-automatic and automatic segmentations.

**Fig 5.** 3D reconstruction of the teeth for the three scans.

**Fig 6.** The alignment of the two scans (Scan 1 and Scan 2) where Scan 1 was used as reference. The grey scale pixel intensity values between 0 and 255 correspond to red, green, and blue values, respectively. Blue is for values closer to 0, green is for intermediate values, and red is for values closer to 255.

**Fig 7.** The alignment of the two reconstructions (Scan 2 and Scan 1) where Scan 1 is taken as reference point cloud. The colors correspond to the distances between the two point clouds. Red is for larger distances, blue is for smaller distances, and green is for medium distances.

**Fig 8.** The alignment of the two reconstructions (Scan 3 and Scan 1) where Scan 1 is taken as reference point cloud. The colors correspond to the distances between the two point clouds. Red is for larger distances, blue is for smaller distances, and green is for medium distances.
notation is similar to figure 7. To eliminate the impact of the outlier points caused by the extra tooth, we excluded the points with distance larger than 500 microns during the statistical analysis. The mean distance and standard deviation between the points from the two scans (at tooth level) are 0.235 and 0.206 mm, respectively. Figure 9 shows the 3D reconstruction at the teeth and gingival levels. Figure 10 shows the aligned reconstructions of the teeth and gingiva from Scan 2 and Scan 1. The mean distance and standard deviation between the points from Scan 2 and Scan 1 are 0.191 and 0.186 mm, respectively.

**Discussion**

The challenge of segmenting the periodontal structures has three main aspects. First, low signal-to-noise ratio, low signal-to-contrast ratio and motion artifact blur the tissue boundaries. Second, different tissues share the same intensity distribution, i.e., the tooth and alveolar bone typically have similar intensity ranges. Also, the echoes from the surface of the gingiva could be strong, which leads to intensity values close to those of bone and tooth. Last, the shape of the structures varies in different sections of the dentition, especially in the transitional area of adjacent teeth. In this area, periodontal structures show significantly different patterns.

The semi-automatic or automatic segmentation of medical US images has been extensively reported in literature, and a review of those applications and techniques was performed by Meiburger et al [22]. However, there are limited reports on US image segmentation in dentistry. Pekam et al [23] used the model-based surface point matching segmentation method to acquire the surface of the prepared artificial teeth model to facilitate prosthetic manufacturing. Xu et al [24] traced the contour in the US tongue image sequence using the active contour method with an automatic re-initialization technique. Chifor et al [25] explored the usage of US in the detection of gingival sulcus with the assistance of active contour based semi-automatic sulcus segmentation in the swine specimen; Nguyen et al [26] segmented alveolar bone in US images using the graph-cut based method with mandibular incisors from piglets, and subsequently adopted a deep learning approach to detect alveolar bone in humans [4].

The major limitation of our semi-automatic segmentation method was the long segmentation time (45 seconds per frame). To overcome this drawback, a deep learning model for object detection and image segmentation was developed. After training, the algorithm was able to automatically detect and segment the three types of tissues. The developed machine learning based segmentation algorithm was a two-order of magnitude faster. It can segment one image in 0.45 second. At the same time, the algorithm demonstrated high accuracy with all tissues identified, and 85% of the segmented area corresponds to that identified by the semi-automated algorithm.

The accuracy of our model is at the same level or above the average of results presented in literature [27-29]. The average accuracy from 25 independent studies published in a recent systematic review, is 88.6% for sensitivity and 93.9% for specificity [30]. However, the reported study included all types of medical imaging modalities (radiography, CT, US) applied to various body regions, while our algorithm used only US, which is known to be more difficult to interpret. Another publication on algorithms for US images of muscle only reported an accuracy of 76% [31]. When the discussion is restricted to the maxillofacial region, the reported deep learning algorithms were applied mostly to radiological and CT imaging. The accuracy of these techniques ranges from 80% [32] to 87% [33]. Rana et al [34] reported an accuracy of about 75% when processing photographic dental images.

The 3D reconstruction discrepancy in the present study compared with other studies is lower. Marotti et al also used CCOSS to calculate the mean distance and reported a reconstruction precision between 12.34 to 46.38 µm [35]. In the Marotti et al [35] study, only one
tooth was examined compared with 4 teeth examined in our study. In regard to the accuracy of CBCT, Tomasi et al reported 0.36 mm for the horizontally positioned mandible, 0.48 mm for the inclined mandible and an overall absolute mean measurement error of 0.40 mm [36], while our study rendered a better accuracy between 0.179 mm and 0.235 mm.

Our paper has certain limitations: the examinations were performed on the same porcine mandible generating probably an overestimated accuracy for the predictions made using artificial intelligence algorithms. Also, it was observed that the prediction underestimated the area of the tissue (refer to Figure 4 for example), most probably due to the low number of training images. Further effort is needed in the future to improve the results using a higher number of highly accurate segmented US periodontal images for training and validation. Despite these limitations the results regarding the reproducibility of the 3D US reconstructions from consecutive scans, using our handheld prototype are promising, qualitatively and quantitatively. The main advantage of the proposed 3D ultrasound reconstruction method is the free-hand architecture of the 3D US scanner prototype, with which one can examine a full maxillary arch.

**Conclusions**

The present feasibility study demonstrated the diagnostic potential of 3D reconstruction using a free-hand 2D US scanner with spatial positioning readings. The use of automated data processing with deep learning algorithms makes the process practical in the clinical environment for the assessment of periodontal soft tissues. Further research should explore reconstruction algorithms with/without interpolation between frames to improve the accuracy and resolution of the reconstruction results. Proper display of the 3D images corresponding to tissue types is also important for the clinical application.

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**Conflict of interest:** none

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