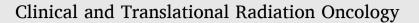
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Design and evaluation of a deep learning-based automatic segmentation of maxillary and mandibular substructures using a 3D U-Net

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ARTICLE INFO ABSTRACT Keywords: Background: Current segmentation approaches for radiation treatment planning in head and neck cancer patients Jaw (HNCP) typically consider the entire mandible as an organ at risk, whereas segmentation of the maxilla remains Dental rehabilitation uncommon. Accurate risk assessment for osteoradionecrosis (ORN) or implant-based dental rehabilitation after Osteoradionecrosis radiation therapy may require a nuanced analysis of dose distribution in specific mandibular and maxillary Radiation therapy segments. Manual segmentation is time-consuming and inconsistent, and there is no definition of jaw Artificial intelligence subsections. Head and neck cancer Materials and methods: The mandible and maxilla were divided into 12 substructures. The model was developed from 82 computed tomography (CT) scans of HNCP and adopts an encoder-decoder three-dimensional (3D) U-Net structure. The efficiency and accuracy of the automated method were compared against manual segmentation on an additional set of 20 independent CT scans. The evaluation metrics used were the Dice similarity coefficient (DSC), 95% Hausdorff distance (HD95), and surface DSC (sDSC). Results: Automated segmentations were performed in a median of 86 s, compared to manual segmentations, which took a median of 53.5 min. The median DSC per substructure ranged from 0.81 to 0.91, and the median HD95 ranged from 1.61 to 4.22. The number of artifacts did not affect these scores. The maxillary substructures showed lower metrics than the mandibular substructures. Conclusions: The jaw substructure segmentation demonstrated high accuracy, time efficiency, and promising results in CT scans with and without metal artifacts. This novel model could provide further investigation into dose relationships with ORN or dental implant failure in normal tissue complication prediction models.

1. Introduction

Head and neck cancer (HNC) is the seventh most prevalent malignancy worldwide [1,2]. Radiation therapy (RT) plays a pivotal role in the treatment of HNC, either as monotherapy, concurrent with chemotherapy, or as adjuvant therapy [3]. Survival rates remain poor at approximately 68,5% after five years, depending on cancer stage and diagnosis [4]. However, studies have shown a better chance of long-term survival for human papillomavirus (HPV)-positive cancers at approximately 80 % after five years [5]. Therefore, it is important to address both the short- and long-term side effects of RT. Despite advancements in radiation therapy techniques, the incidence of osteor-adionecrosis (ORN) remains high, with reported rates of up to 10 % [6,7].

To prevent odontogenic complications during and after RT, it is necessary to extract loose, periodontally infected, and severely decayed teeth prior to therapy, as extractions are particularly susceptible to the risk of ORN [8–10]. Complications can lead to the discontinuation of RT,

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which can significantly reduce the probability of cancer cure or contribute to long-term morbidity [11]. Extractions can result in partial or complete edentulism, and health-related quality of life (HRQoL) studies have shown that patients often experience significant difficulties with eating, swallowing, and speaking [2,12,13]. Dental implant (DI)-supported dentures are the preferred solution for restoring oropharyngeal function [14–17]. This approach requires strategically placed dental implants in the mandible and maxilla for secure attachment, potentially necessitating large bone exposures or bone augmentation [18]. These post-radiotherapeutic dentoalveolar procedures also pose a risk for ORN, with the administered radiation dose being a contributing factor that can even affect DI survival [16,19–24].

Historically, the maximum dose for the entire mandible was limited to 70 Gy for a volume of 0.1 cm^3 . Recent recommendations suggest using volume constraints. For instance, it is recommended to limit the volume receiving 44 Gy to no more than 42 % or 58 Gy to no more than 25 % [25].

Nevertheless, ORN appears to be more prevalent in the premolar, molar, and retromolar regions than in other regions of the mandible, indicating that the jaws should be considered in a more differentiated approach [6].

Dental implants are considered safe for patients who receive a mean radiation dose of less than 38 Gy. In contrast, a high failure rate has been observed when the radiation dose exceeds 50 Gy, even if the implantation occurs over two years post-RT [16,19-23].

Further research is needed to determine whether a defined dose reduction in specific areas of the jaw will reduce ORN prevalence or increase DI survival. Additionally, a better understanding of jaw tolerances is necessary [25], which might enhance risk management.

Modern RT techniques, such as intensity-modulated RT (IMRT) or volumetric intensity-modulated arc therapy (VMAT), have increasingly replaced three-dimensional (3D) conventional planning, leading to more homogeneous and conformal irradiation and making sparing surrounding risk structures feasible [26,27]. Manual segmentation is necessary to use the maxilla and mandible as organs at risk (OAR) in RT planning. However, this process is resource-intensive, time-consuming, and repetitive. Challenges arise due to high-attenuation materials causing artifacts, variability in the maxillary sinus, patient-specific variations, and the complexity added by the proximity of other bony structures and the overlap of upper and lower teeth in adjacent CT slices [28-30]. Recent technological advancements have introduced automated medical image segmentation to alleviate clinical workload. Several studies have proposed semi-automated or fully automated methods for mandible segmentation in CT or cone beam CT scans [28-35]. These methods serve various purposes, such as OAR delineation, diagnosis, or maxillofacial surgery planning. Among the segmentation techniques, those based on deep learning (DL) architectures, such as U-Net, have shown superior performance in addressing mandible segmentation challenges. Ibragimov and Xing demonstrated the pioneering application of DL techniques using convolutional neural networks (CNN) for segmenting OARs in head and neck CT scans [34]. However, previous research has primarily focused on the mandible, with limited exploration of comprehensive techniques for automatically segmenting the maxilla. Moreover, studies have not yet attempted to segment the jaw into substructures to spare specific areas or illustrate dose distributions. This could further investigate dose relationships with ORN or implant failure in normal tissue complication prediction (NTCP) models [36].

To address this gap, this study presents a new model that automatically segments the maxilla and mandible into twelve anatomically based substructures using a 3D encoder-decoder U-Net [37] architecture.

2. Methods and materials

2.1. Study protocol

Before participating in the study, all patients involved provided informed consent. The study's protocol received approval from the Ethics Committee of Charité-Universitätsmedizin in Berlin, Germany, under the reference number EA2/155/21. The research was conducted following the principles of the Declaration of Helsinki.

2.2. Patient selection

The study used 82 CT scans to train and validate the model. The training set consisted of 64 CT scans, of which 53 were from Charité-Universitätsmedizin Berlin, Germany, and 11 were from MVision AI Oy, Helsinki, Finland, in compliance with the General Data Protection Regulation. The validation set included 18 scans, 15 from Charité and three from MVision AI Oy. After the development phase, the model underwent testing on an additional set of 20 independent CT scans obtained exclusively from Charité. The scans from Charité were from patients diagnosed with head and neck cancer and undergoing (chemo) radiation therapy. The group comprised 72 % males and 28 % females. with an average age of 66.8 years. The scans were collected between January 2013 and October 2020 using Siemens Somatom Sensation Open scanners. The median slice thickness was 2 mm (range 1–3 mm), and metal artifact reduction was not applied. Scans provided by MVision included five males, two females, and four individuals of unspecified gender, with specific age information available for three individuals: 42, 50, and 82 years. Siemens Sensation Open and Siemens Somatom Definition AS scanners were used for imaging, with a slice thickness ranging from 2 to 3 mm. Imaging data was missing for three cases. All scans had a slice size of 512 x 512 pixels, except for one which had a size of 256 x 256 pixels. CT scans with osseous destruction of the jawbone and HNC patients with a tumor site in the oral cavity were excluded.

2.3. Jaw segmentation

Dental reference structures for prosthetic restorations were used to determine segment boundaries in the mandible and maxilla. Therefore, the mandible (lower jaw, LJ) was divided into eight segments and the maxilla (upper jaw, UJ) into four segments. In cases where teeth were present, the dental crown was omitted, and only the bony structure was delineated. An interdisciplinary team of radiation oncologists and dental professionals clearly defined the criteria for delineating each segment. A summary of the segment definition is provided in Table 1. An example of the manual segmentation of the mandible and maxilla is shown graphically in representative CT slices in Fig. 1. Segments were delineated using EclipseTM version 15.5 (Varian Medical Systems, Inc., Palo Alto, CA, USA) and the MVision AI online annotation platform, version 3.1. Dental status, including metal-related artifacts, was analyzed in each segment.

2.4. Model training and validation

The training dataset underwent extensive data augmentation to enhance model generalization and robustness. Preprocessing included a random selection of three consecutive slices from CT images, followed by cropping and padding to achieve a standardized axial patch size of 512 x 512 pixels. Intensity transformations such as random gamma transform, gamma inversion, additive and multiplicative intensity transform, Gaussian noise, Gaussian blur, and sharpening were applied with specified probabilities. Additionally, spatial transforms were employed, including random deformable transforms and scaling. Finally, intensity values were normalized with specified mean and standard deviation parameters and clamped within a predefined range.

The segmentation model adopts an encoder-decoder U-Net structure,

Table 1

Segments and Boundary Definitions of the 12 segments of the Jaw Model.

	Boundaries and Definition							
Structure	Cranial	Caudal	Dorsal	Lateral	Medial	Anterior		
Upper jaw front left and right (UJFL/UJFR)	Base of the anterior nasal spine	Basilar edge of the maxilla	Dorsal edge of the maxilla	Plane tangent to the lateral edge of the nose ("canine line")	Center line through both incisive foramen	Anterior edge of the maxilla		
Upper jaw molar left and right (UJML/ UJMR)	Base of the anterior nasal spine/base of the maxillary sinus		Connecting line between both greater palatine foramen	Lateral edge of the maxilla	Medial edge of the maxilla	Plane tangent to the lateral edge of the nose ("canine line")		
Lower Jaw front left and right (LJFL/LJFR) Lower Jaw molar left and right (LJML/	Apical edge of the mandible (without teeth)	Basilar edge of the mandible	Plane tangent to the mental foramen Frontal plane: Connecting line between the body of the mandible and the ramus of	Lateral edge of the mandible	Medial edge of the mandible	Anterior edge of the mandible / mental protuberance Plane tangent to the mental foramen		
LJMR) Lower Jaw ramus left and right (LJRL/ LJRR) Lower Jaw condyle left and right (LJCL/LJCR)	Anterior part: Apical edge ramus of the posterior part of the mandible: Connecting line to LJ condyle left/right Cranial surface of the condylar head	Connecting line to LJ ramus left/ right	the mandible Dorsal edge of the mandible			Frontal plane: Connecting line between the body of the mandible and the ramus of the mandible Mandibular notch		

with the encoder utilizing a Residual Neural Network (ResNet)-type [38] backbone pre-trained on ImageNet [39] to extract hierarchical features from input CT images. These features are symmetrically passed with skip connections to the decoder, comprising multiple Dense Convolutional Network (DenseNet) [40] blocks. Two Upsampling Convolutional (UpConv) [41] layers with 2x upsampling are then employed, followed by a 3D convolutional (Conv 3D) layer to generate final segmentations, with the same spatial size as the original input CT volume. The full architecture is shown in Fig. 2. The DenseNet decoder block is shown in Fig. 3 (a). The output of the previous layer (x_1) is upsampled using bilinear interpolation to match the spatial size of the input provided by the skip connection (x₂). Then, the two tensors are concatenated and passed through a series of Batch Normalization (BN) [42], Rectified Linear Unit (ReLU), and two-dimensional convolutional (Conv 2D) layers. Finally, the output is concatenated to the original input of this sequence of layers. The UpConv 2x layer is shown in Fig. 3 (b). After the input is passed through a BN, a ReLU, and a Conv 2D layer, it is upsampled using bilinear interpolation to have twice the spatial size compared to its original size.

The training procedure employed the sum of Dice loss and weighted cross-entropy loss to optimize the model's performance. The segmentation model utilized the Adam [43] optimizer with a learning rate of 3e-4, momentum of 0.9, and weight decay of 3e-5. The training process spanned 123 epochs, with each epoch comprising 250 iterations. The model was trained with a dual NVIDIA RTX 3090 GPU system (24 GB VRAM each), 64 GB of RAM, and an AMD Ryzen 9 5900X 12-Core CPU. The total training time was approximately 6.5 h.

A ReduceLROnPlateau learning rate scheduler was implemented with a threshold of 1e-2, dynamically adjusting the learning rate during training based on the model's performance plateau.

The cloud-based system uses pseudonymized DICOM data for automated segmentation and is connected to the clinic's RT planning system. A team of six annotators delineated the training and validation datasets, all with backgrounds in medical radiation oncology and dentistry and specific expertise in head and neck radiological anatomy. The bony structures were easily distinguishable from the surrounding tissue, and minor corrections were needed, as judged by two experienced head and neck radiation oncologists.

2.5. Quantitative evaluation

An independent set of 20 CTs was selected from the Charité-Universitätsmedizin Berlin dataset to evaluate the performance of the model (testing). Manual and automated segmentation were performed on each CT. Both segmentations were compared using quantitative scoring with the Dice Similarity Coefficient (DSC), 95 % Hausdorff Distance (HD95), and the surface DSC (sDSC). Manual and automated segmentation times were measured. The test dataset was delineated through manual segmentation by one annotator with a background in dentistry.

2.6. Statistics

The patient cohort was divided into a median split based on the number of segment-related artifacts. Patients with either no metal artifacts or a maximum of one artifact in a segment were compared to patients with more than one artifact in a segment. The number of segments with artifacts was non-parametrically correlated with DSC and HD95 using Spearman's correlation coefficient. In addition, the subgroups created by the median splitting in the patient sample were compared using non-parametric Mann-Whitney U tests. A p value ≤ 0.05 was considered statistically significant. MATLAB® R2021a (The MathWorks, Inc., Natick, MA, USA), IBM SPSS Statistics for Windows, version 28.0 (IBM Corp., Armonk, NY, USA), and Microsoft Excel 365 (Microsoft Corporation, Redmond, WA, USA) were used for statistical analysis.

3. Results

3.1. Jaw segmentation

The 3D Jaw Model created after segmentation and color subdivision is shown in Fig. 4.

The dental status of the training and validation dataset is shown in Table 2 and varied between full dentition, partial dentition (one to 15 teeth), and edentulous patients. The metal-related artifacts in the mandible and maxilla were listed.

3.2. Quantitative evaluation

The median number of CT slices in the test dataset of 20 patients was

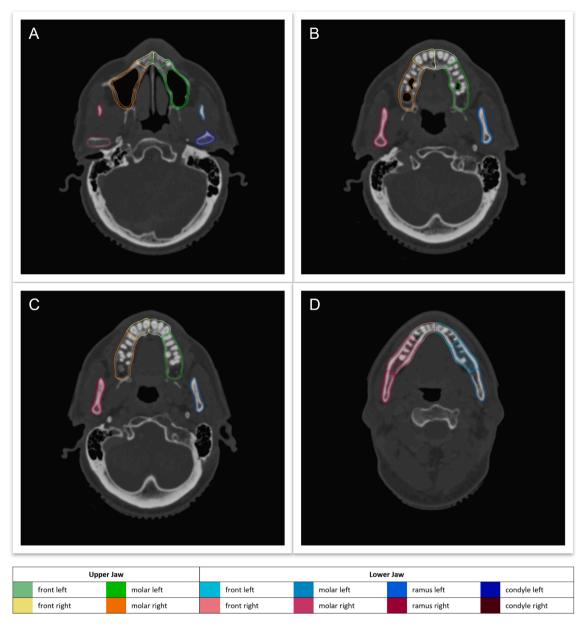


Fig. 1. Axial cross-sectional CT images of the jaw illustrating manual segmentation. (A) shows the maxilla with the maxillary sinus intentionally left unsegmented; (B) and (C) provide additional axial views of the segmented maxilla; (D) shows the axial view of the manually segmented mandible.

191 (range 108–314), and the median slice thickness was 2 mm (range 1–2.0 mm). The automated segmentations were generated in a median of 86 s (range 39–150 s). Conversely, the manual segmentations required a median time of 53.5 min (range 36–75 min). The comparison between manual and automated segmentation showed a maximum median HD95 of 4.22 mm for the LJML segment, while the LJCL segment showed a minimum median HD95 of 1.61 mm. The UJML segment had the widest range for HD95 at 11.01 mm, while the LJFR segment had the narrowest range at 4.57 mm. The maximum and minimum median DSC values were 0.91 for LJRR and 0.81 for UJML. The highest sDSC value (0.98) was observed for LJFR, while the lowest (0.87) was found for UJFR. A comprehensive summary of the results is presented in Table 3, Fig. 5, and Fig. 6.

According to the Mann-Whitney U test, the subgroup analysis using a median split based on the number of artifact-affected segments showed no significant differences between the subgroups in quantitative scores (p = 0.579 for DSC and HD95). In addition, non-parametric Spearman correlation analysis revealed no significant relationships between the

number of artifact-affected segments and the median DSC and HD95 (DSC: p = 0.628; HD95: p = 0.780).

4. Discussion

This study presents a novel 3D encoder-decoder U-Net-based model that enables the segmentation of the jaw into 12 substructures.

Automated segmentation has emerged as a pivotal tool in radiation planning, enhancing accuracy, reproducibility, and efficiency, thereby reducing clinical workload and standardizing treatments [28–35,44]. Segmenting the entire mandible, a feature prevalent in many autosegmentation tools, primarily aims to minimize the risk of ORN. However, the variable tumor locations and the associated diverse dose distributions within the entire mandible expose the limitations of current segmentation in dental rehabilitation post-RT [6]. Furthermore, segmentation of the maxilla is not always performed. In an independent test dataset, the model proved to be effective for segmentation. Manual segmentation of the jaw substructures in our study was time-consuming,

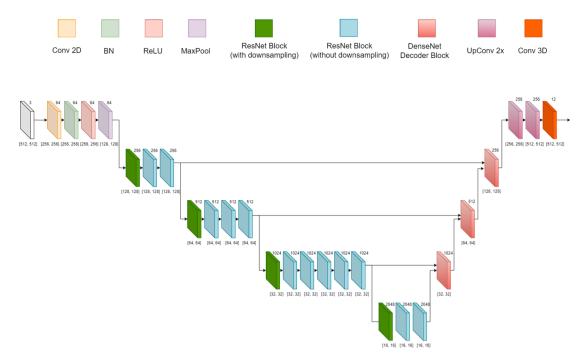


Fig. 2. Schematic representation of the U-Net architecture composed of a ResNet-based encoder and DenseNet decoder for feature extraction and image segmentation. The numbers above each layer denote the output feature size of the respective layer, while the numbers below the layers denote the spatial size of their output. Abbreviations: BN = Batch Normalization, ReLU = Rectified Linear Unit, Conv 2D = Two-Dimensional Convolutional Layer, Conv 3D = Three-Dimensional Convolutional Layer, UpConv = Upsampling Convolutional Layer, MaxPool = Maximum Pooling, ResNet = Residual Neural Network, DensNet = Dense Convolutional Network.

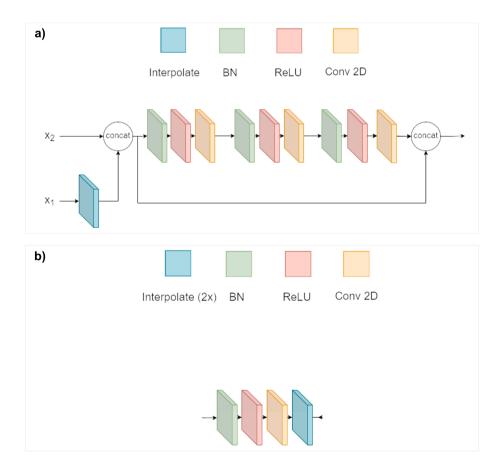


Fig. 3. Detailed illustrations of the decoder components within the U-Net architecture, featuring a) the DenseNet block and b) the UpConv 2x layer. Abbreviations: BN = Batch Normalization, ReLU = Rectified Linear Unit, Conv 2D = Two-Dimensional Convolutional Layer, Conv 3D = Three-Dimensional Convolutional Layer.

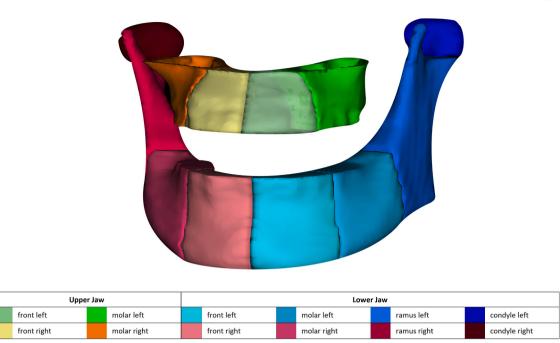


Fig. 4. Jaw Model – Segmentation of the Mandible and Maxilla into 12 substructures. The color-coded legend beneath the model delineates each segment. Upper Jaw (UJ) from left to right: UJ molar right, UJ front right, UJ front left, UJ molar left. Lower Jaw (LJ) from left to right: LJ condyle right, LJ ramus right, LJ molar right, LJ front right, LJ front left, LJ ramus left, LJ condyle left.

Table 2 Dental Status of the Training, Validation and Test Dataset.

	Upper Jaw	Lower Jaw
Training dataset ($n = 64$)		
Full dentition	14	13
Partially edentulous	18	22
Edentulous	32	29
Metal-related artifacts	20	25
Validation dataset (n = 18)		
Full dentition	1	1
Partially edentulous	5	9
Edentulous	12	8
Metal-related artifacts	4	7
Test dataset ($n = 20$)		
Full dentition	12	13
Partially edentulous	7	6
Edentulous	1	1
Metal-related artifacts	13	10

taking approximately 53 min. Artifacts and complex structures such as the maxillary sinus and precise adherence to substructure boundaries contributed to more complex and time-consuming delineation. In contrast, the automated segmentation process in our study took an average of 86 s, surpassing Ibragimov's initial work on OAR segmentation in the head and neck region using CNN networks, which took approximately 4 min [34]. Morita et al., who developed a twodimensional U-net architecture for facial bone segmentation, were able to reduce the required time to an average of 10 s per region (e.g., maxilla and mandible) [35]. Our longer output time could be explained by the fact that our model is additionally divided into subsections.

The evaluation of the automated segmentation in this study involved multiple metrics, using manual segmentation as the ground truth. An independent test dataset analysis showed a median DSC of 0.88, a median HD95 of 3.00 mm, and a median sDSC of 0.94 % per patient across all substructures. These results demonstrate robust performance despite variations in patient characteristics such as artifact count, dental status, and bone volume. However, this study represents the first automated segmentation of maxillary and mandibular substructures, making direct

Table 3

Quantitative Evaluation comparing the manual and automated segmentation of
the Jaw Model in the test dataset per substructure.

	Performance metric							
	DSC (median, range)	HD95 [mm] (median, range)	sDSC [%] (median, range)					
Whole UJ	0.82	3.31	0.89					
Whole LJ	0.89	2.80	0.96					
Substructures								
UJFL	0.82 (0.67-0.89)	3.61 (2.29-7.20)	0.87 (0.70-0.98)					
UJFR	0.81 (0.66-0.88)	3.16 (2.05-7.24)	0.88 (0.70-0.99)					
UJML	0.81 (0.58-0.90)	3.46 (1.43-12.43)	0.90 (0.62-0.99)					
UJMR	0.82 (0.52-0.89)	2.87 (2.09-12.18)	0.91 (0.62-0.99)					
LJFL	0.88 (0.76-0.94)	2.99 (1.56-6.86)	0.97 (0.75–1.00)					
LJFR	0.88 (0.81-0.92)	2.61 (1.93-6.50)	0.98 (0.80-1.00)					
LJML	0.88 (0.80-0.98)	4.22 (1.12-8.15)	0.89 (0.81-1.00)					
LJMR	0.90 (0.81-0.94)	3.22 (1.76-9.20)	0.93 (0.81-0.99)					
LJRL	0.88 (0.79-0.98)	3.02 (0.40-6.40)	0.95 (0.87-1.00)					
LJRR	0.91 (0.79-0.95)	2.41 (1.08-10.01)	0.97 (0.85-1.00)					
LJCL	0.90 (0.70-0.95)	1.61 (1.01-6.64)	0.96 (0.75–1.00)					
LJCR	0.89 (0.63–0.94)	1.62 (1.05–7.46)	0.96 (0.73–1.00)					

Legend: DSC = Dice surface coefficient, HD95 = 95 % Hausdorff Distance, sDSC = surface DSC. UJFL = upper jaw front left, UJFR = upper jaw front right, UJML = upper jaw molar left, UJMR = upper jaw molar right, LJFL = lower jaw front left, LJFR = lower jaw front right, LJML = lower jaw molar left, LJMR = lower jaw molar right.

comparisons with previous studies difficult.

Furthermore, there is no uniform definition of which metric levels are sufficient for clinical use. The multi-atlas segmentation strategy, a traditional method, achieved a DSC of 0.86 for the entire mandible [33]. However, these conventional methods have been surpassed by CNNs due to their inability to handle high noise or metal artifacts and the need for expert analysis, showing poor individualization for mandibular segmentation cases [45].

Ibragimov and Xing observed a median DSC of 0.89 using CNN [34]. The DSC could be improved by combining methods, as shown by Tong et al. [46]. They combined a fully convolutional neural network (FCNN) with a shape representation model (SRM), achieving a DSC of 0.92 for

	Substructure											
Patient	UJFL	UJFR	UJML	UJMR	LJFL	LJFR	LJML	LJMR	LJRL	LJRR	LJCL	LJCR
1	0,79	0,84	0,81	0,81	0,89	0,88	0,91	0,85	0,89	0,80	0,88	0,87
2	0,89	0,82	0,58	0,52	0,81	0,83	0,81	0,89	0,84	0,86	0,82	0,88
3	0,82	0,86	0,78	0,75	0,90	0,91	0,87	0,90	0,84	0,89	0,79	0,84
4	0,83	0,80	0,59	0,73	0,82	0,81	0,84	0,91	0,85	0,93	0,86	0,93
5	0,80	0,80	0,85	0,80	0,89	0,91	0,88	0,94	0,88	0,94	0,88	0,90
6	0,80	0,84	0,87	0,89	0,89	0,88	0,89	0,92	0,88	0,92	0,94	0,89
7	0,82	0,76	0,80	0,86	0,91	0,88	0,90	0,86	0,89	0,88	0,85	0,78
8	0,82	0,82	0,90	0,89	0,89	0,87	0,91	0,90	0,93	0,91	0,93	0,90
9	0,87	0,85	0,87	0,88	0,87	0,87	0,92	0,91	0,90	0,91	0,92	0,89
10	0,67	0,68	0,80	0,86	0,83	0,87	0,85	0,89	0,86	0,91	0,89	0,94
11	0,68	0,80	0,70	0,76	0,94	0,92	0,89	0,84	0,89	0,85	0,85	0,75
12	0,84	0,88	0,81	0,82	0,86	0,87	0,86	0,90	0,87	0,91	0,94	0,94
13	0,81	0,81	0,86	0,86	0,88	0,87	0,90	0,90	0,90	0,93	0,90	0,88
14	0,82	0,86	0,88	0,85	0,90	0,92	0,88	0,94	0,85	0,92	0,89	0,94
15	0,83	0,78	0,80	0,83	0,87	0,89	0,80	0,81	0,79	0,79	0,70	0,63
16	0,75	0,81	0,83	0,82	0,93	0,91	0,91	0,89	0,88	0,85	0,91	0,87
17	0,71	0,66	0,80	0,82	0,87	0,86	0,98	0,93	0,98	0,95	0,92	0,91
18	0,86	0,81	0,87	0,82	0,76	0,82	0,83	0,87	0,89	0,92	0,95	0,94
19	0,84	0,88	0,80	0,88	0,89	0,91	0,87	0,94	0,84	0,88	0,92	0,90
20	0,73	0,70	0,64	0,68	0,88	0,89	0,88	0,90	0,88	0,93	0,93	0,89
L												
Median	0,82	0,81	0,81	0,82	0,88	0,88	0,88	0,90	0,88	0,91	0,90	0,89

Fig. 5. Heatmap Visualization of DSC across all substructures and cases.

the FCNN alone and 0.94 with the SRM-constrained FCNN. Liu et al. evaluated 19 models from 18 studies for automated segmentation of the entire mandible and found a pooled DSC of 0.92 [47]. Previous studies have reported HD95 values ranging from 1.98 to 2.83 [48,49]. The study found that there were performance differences between the sub-structures, with the maxillary substructures generally having lower metrics than the mandible, as shown in Table 3, Fig. 5 and Fig. 6.

In this study, the median DSC for the entire maxilla was 0.82, compared to 0.89 for the mandible when the substructures were combined. Furthermore, the maxilla's HD95 values were worse than those for the mandible (3.31 vs. 2.8). The maxillary molar substructures showed higher standard deviations in performance metrics, indicating

variability in segmentation accuracy due to the maxilla's complex anatomy and a higher incidence of artifacts (65 %). However, artifacts did not significantly affect the overall performance metrics, which is consistent with the results presented by Ilesan et al. [50]. They reported a similar DSC for automated mandibular bone segmentation with and without artifacts.

Morita et al. also had a worse DSC for segmentation of the entire maxilla compared to the entire mandible (0.91 vs. 0.98) [35]. Training and validating the model with additional cases could increase the consistency of maxillary segmentation performance to improve its robustness.

Overall, the metrics' performance is inferior to that of the compared

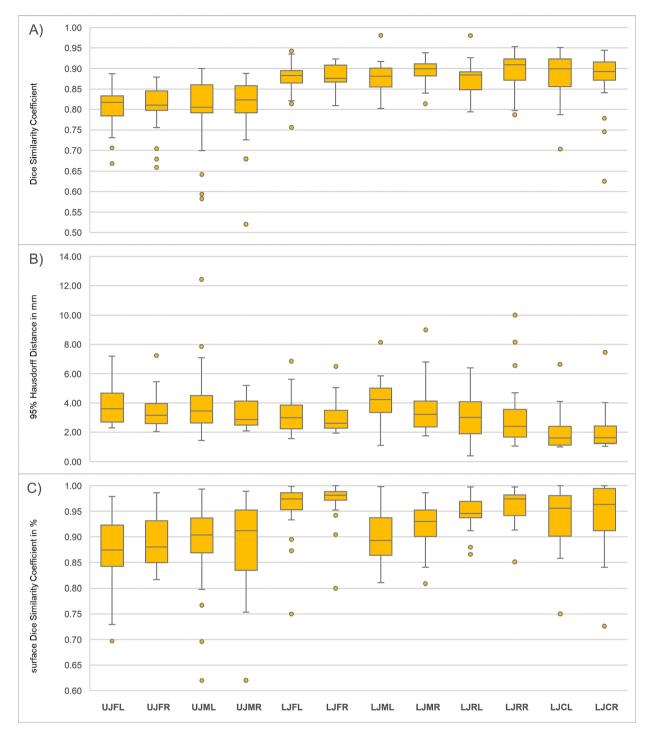


Fig. 6. Box and whisker plots comparing the manual and automated segmentation of the Jaw Model in the test dataset. A) Dice Similarity Coefficient, B) 95 % Hausdorff Distance, C) surface Dice Similarity Coefficient. The median is shown as a horizontal line within each box, while yellow dots represent outliers (exceeding 1.5 times the interquartile range of the upper and lower quartiles). Abbreviations: UJFL = upper jaw front left, UJFR = upper jaw front right, UJML = upper jaw molar left, UJMR = upper jaw molar right, LJFL = lower jaw front left, LJFR = lower jaw molar right, LJRL = lower jaw molar right, LJCL = lower jaw condyle left, LJCR = lower jaw condyle right. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of this article.)

studies. It should be noted that the division into substructures within the same density of tissue is subject to increased variability since the delineation can be more difficult.

In this study, the boundaries are always bone-to-bone structures with equal Hounsfield units compared to bone-to-tissue boundaries. Further investigation is required to determine whether the present results would be better if the maxilla and mandible were considered a single segment. One limitation of this study is that the Jaw Model mainly uses scans from a single institution. This may limit its ability to adapt to CT scans with different imaging characteristics, potentially affecting the evaluation metrics. Additionally, the method was only applied to intact jawbones with standard dental artifacts, excluding those with complex artifacts such as reconstruction plates that could compromise segmentation integrity. It is important to note that the absence of analysis of interobserver and intraobserver variability is due to the precise segmentation boundaries requiring minimal adjustments. The evaluation of the model on only 20 CT scans highlights the need for further improvement and broader validation to enhance its robustness and clarify its limitations.

5. Conclusions

The results of this study demonstrate the feasibility of dividing the maxilla and mandible into subsections using the 3D U-Net architecture. The model's ability to generate a larger, more coherent data set promises significant contributions to the research field. This will allow further investigations into ORN development, more detailed dose distributions within the jaws, and, therefore, the feasibility of sparing segments or even improving dental rehabilitation.

CRediT authorship contribution statement

L. Melerowitz: Conceptualization, Writing – original draft, Investigation, Methodology, Validation, Visualization, Formal analysis, Data curation. S. Sreenivasa: Conceptualization, Methodology, Writing – review & editing, Formal analysis. M. Nachbar: Writing – review & editing, Methodology, Formal analysis. A. Stsefanenka: Writing – review & editing, Data curation. M. Beck: Writing – review & editing. C. Senger: Writing – review & editing. N. Predescu: Methodology, Software, Writing – review & editing. S. Ullah Akram: Methodology, Software. V. Budach: Writing – review & editing. D. Zips: Writing – review & editing. M. Heiland: Writing – review & editing. S. Nahles: Writing – review & editing, Supervision, Conceptualization, Methodology, Project administration. C. Stromberger: Conceptualization, Methodology, Writing – review & editing, Supervision, Project administration.

Declaration of competing interest

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: The authors disclose institutional collaboration with MVision AI, which was responsible for the technical development of the presented model. MVision AI did not participate in the analysis or evaluation of the test dataset.

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