

GEPAR3D: Geometry Prior-Assisted Learning for 3D Tooth Segmentation

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Abstract. Tooth segmentation in Cone-Beam Computed Tomography (CBCT) remains challenging, especially for fine structures like root apices, which is critical for assessing root resorption in orthodontics. We introduce **GEPAR3D**, a novel approach that unifies instance detection and multi-class segmentation into a single step tailored to improve root segmentation. Our method integrates a Statistical Shape Model of dentition as a geometric prior, capturing anatomical context and morphological consistency without enforcing restrictive adjacency constraints. We leverage a deep watershed method, modeling each tooth as a continuous 3D energy basin encoding voxel distances to boundaries. This instance-aware representation ensures accurate segmentation of narrow, complex root apices. Trained on publicly available CBCT scans from a single center, our method is evaluated on external test sets from two in-house and two public medical centers. **GEPAR3D** achieves the highest overall segmentation performance, averaging a Dice Similarity Coefficient (DSC) of 95.0% (+2.8% over the second-best method) and increasing recall to 95.2% (+9.5%) across all test sets. Qualitative analyses demonstrated substantial improvements in root segmentation quality, indicating significant potential for more accurate root resorption assessment and enhanced clinical decision-making in orthodontics. We provide the implementation and dataset at github.com/tomek1911/GEPAR3D.

Keywords: Tooth segmentation · Geometry prior · Root resorption

1 Introduction

CBCT is essential in digital dentistry, yet manual tooth segmentation remains labor-intensive and inconsistent [32]. Automated methods support treatment planning and diagnostics [14], but delineating tooth roots remains challenging

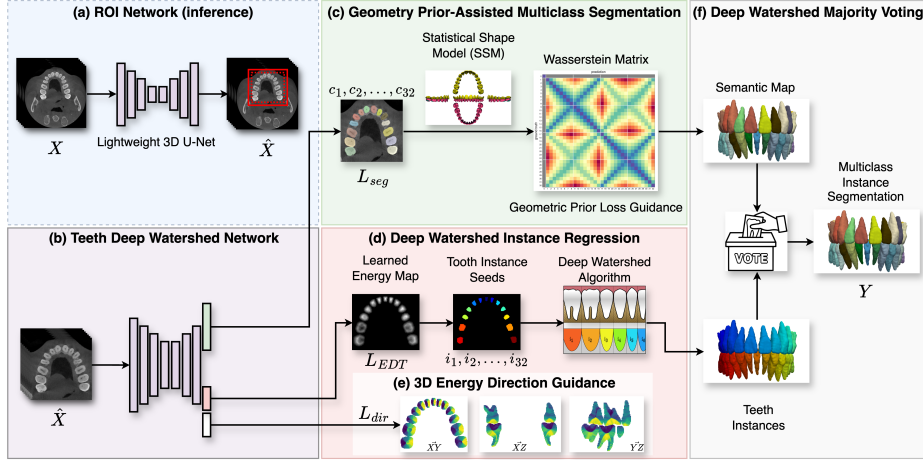


Fig. 1. An overview of GEPAR3D, which unifies instance detection and multi-class segmentation for precise tooth root segmentation. **(a)** Crops the region of interest (ROI) during inference. **(b)** Simultaneously performs multi-class segmentation and instance regression. **(c)** Regularizes segmentation loss L_{seg} with a geometric prior from an SSM of normal dentition [15]. **(d)** Uses instance regression task L_{EDT} to generate energy maps for the Deep Watershed Algorithm. **(e)** Captures complex root apex geometries via Energy Direction loss L_{dir} . Finally, **(f)** assigns each detected instance a class via majority voting based on segmentation outputs.

due to their intricate morphology and small size. Accurate segmentation is particularly important for assessing root resorption [26], a pathological loss of dentin and cementum often caused by orthodontic tooth movement, which can weaken stability and, in severe cases, increase the risk of tooth loss. Tooth geometry exhibits universal patterns, with teeth arranged in two arches and four quadrants, and although individual variability exists, key anatomical features remain stable [17]. Demonstrating this structural consistency, the upper molars typically have three roots, while the lower ones usually have two [31]. Leveraging inherent geometric priors may provide valuable guidance for enhancing the accuracy and robustness of automated segmentation.

Tooth segmentation has evolved from heuristic, hand-crafted methods to deep learning [24]. Early deep learning approaches rely on voxel-wise classification and overlook anatomical structure and inter-class relationships, both essential for capturing detailed tooth morphology, especially root regions [8, 18, 4, 28]. To reduce computational costs, many methods use multi-step coarse-to-fine pipelines that isolate individual teeth in bounding boxes [5, 9, 13, 16, 29]. Such pipelines tend to accumulate errors and disconnect each tooth from its broader anatomical context, impairing root apex segmentation. Recent methods integrate prior anatomical knowledge to guide segmentation. ToothSeg [9] enforces shape consistency via tooth skeletons but relies on manual thresholds and post-processing, limiting generalization. Other methods incorporate spatial relationships via ad-

jacency constraints; SGANET [19] uses rigid graph convolutions to enforce local consistency of immediate neighbors, while TSG-GCN [21] learns dynamic adjacency but remains vulnerable to data bias since its geometric priors derive solely from training labels. These constraints highlight the need for a method that integrates anatomical structure and inter-class dependencies to achieve a coherent, context-rich representation for accurate root segmentation.

To achieve that, we propose **GEometric Prior-Assisted LeaRning for 3D** (GEPAR3D), a novel approach that unifies instance detection and multi-class segmentation in a single step. Our method integrates a Statistical Shape Model (SSM) [15] of dentition as a geometric prior, based on inter-teeth distances, to embed anatomical context and morphological consistency into the learning process. Furthermore, we leverage a deep watershed method and model each tooth as a continuous 3D energy basin, encoding voxel distances to boundaries, and predicting directional gradients to capture subtle variations at root apices. Trained on publicly available CBCT scans from a single center and evaluated on external test sets from four medical centers (two in-house and two public), GEPAR3D demonstrates robust generalization across diverse patient demographics. Our method outperforms five state-of-the-art methods, achieving the highest segmentation performance with an average DSC of 95.0% (+2.8%) and RC of 95.2% (+9.5%) across all test sets, offering new prospects for reliable root resorption assessment and improved clinical decision-making.

2 Methodology

Fig. 1 overviews GEPAR3D, an encoder-decoder model with dual decoders for multi-class segmentation and instance regression. The segmentation branch classifies 32 tooth categories with SSM-based regularization, while the regression branch models instances as energy basins guided by energy descent. Each detected instance receives class votes from the multi-class segmentation branch, and the final class assignment is determined through majority voting.

Geometric prior. To enhance root segmentation in CBCT scans, we integrate an SSM [15] as a geometric prior. This 3D atlas of normal dentition, built from representative Korean individuals (47 males, 37 females), guides segmentation via extracted inter-tooth distances; see [15] for more details on demographics. To capture statistical tooth positions, we represent each tooth’s geometric center as $G_i = (x_i, y_i)$ in a normalized coordinate system, where i denotes the tooth index. Each quadrant Q_k , where $k \in \{1, 2, 3, 4\}$, contains 8 teeth, defined as $T_{Q_k} = \{G_{k1}, G_{k2}, \dots, G_{k8}\}$. The statistical inter-tooth Euclidean distances $D_{ij}^{(k)}$ within a quadrant are computed as: $D_{ij}^{(k)} = \sqrt{(x_i - x_j)^2 + (y_i - y_j)^2}$, with $G_i, G_j \in T_{Q_k}$. The origin O_{xy} of the normalized system is set at the midpoint of the maxillary and mandibular central incisors, whose geometric centers define: $O = \frac{1}{4}(G_{11} + G_{21} + G_{31} + G_{41})$. Since $T_{k,8}$ is absent from the SSM due to rarity, its geometric center G_8 is interpolated from G_6 and G_7 . Finally, statistical inter-tooth distances D_{ij} for each quadrant Q_k are obtained by averaging male and female dentition models, forming the

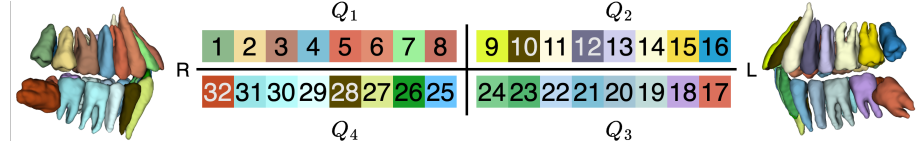


Fig. 2. Quadrants Q_k layout with 32-tooth Universal Numbering System [1].

intra-quadrant distance matrix $\mathbf{D}_{Q_k} = [D_{ij}^{(k)}]$, where D_{ij} encodes pairwise Euclidean distances between teeth within Q_k .

Geometry Prior-Assisted Learning. To ensure semantically meaningful predictions, we adapt a prior-regularized DSC for multi-class segmentation, the Generalized Wasserstein Dice Loss (GWDL) [10], by replacing its original empirical dissimilarity penalties with statistical inter-tooth distances as a geometric prior, yielding the Geometric Wasserstein Dice Loss (GeoWDL). Wasserstein Distance (WD), derived from Optimal Transport (OT) [10], quantifies the minimal cost of transforming one probability distribution into another. When enriched with geometric priors, it enables structured penalization of segmentation errors based on spatial and morphological relationships. Specifically, higher penalties are assigned to errors between distant teeth within the same quadrant, reflecting geometric prior, while semantically weighted adjustments assign lower penalties to misclassifications among morphologically similar teeth within the same arch and higher penalties to confusions between the structurally distinct upper and lower arches. We introduce penalty modifiers $p_{Q_{k_i}Q_{k_j}}$ to weight geometry-based penalties, as matrix $\mathbf{Q}_{q_{ij}}$ ($i, j \in k$) to penalize confusions between quadrants Q_k (Fig. 2): within dental arch ($p_{Q_1Q_2} = 0.1$), between arches ($p_{Q_1Q_4} = 0.2$) and diagonally ($p_{Q_1Q_3} = 0.3$), given as:

$$\mathbf{Q}_{q_{ij}} = \begin{bmatrix} 0 & q_{12} & q_{13} & q_{14} \\ q_{21} & 0 & q_{23} & q_{24} \\ q_{31} & q_{32} & 0 & q_{34} \\ q_{41} & q_{42} & q_{43} & 0 \end{bmatrix} = \begin{bmatrix} 0 & 0.1 & 0.3 & 0.2 \\ 0.1 & 0 & 0.2 & 0.3 \\ 0.3 & 0.2 & 0 & 0.1 \\ 0.2 & 0.3 & 0.1 & 0 \end{bmatrix}. \quad (1)$$

To obtain the penalty matrix \mathbf{M}_{geo} (hereafter M for brevity), we arrange \mathbf{D}_{Q_k} and apply penalty modifiers $\mathbf{Q}_{q_{ij}}$. First, we define the helper matrix $\mathbf{P}_{mn,ij} = q_{ij}J_{mn}$, where \mathbf{J}_{mn} is an 8×8 matrix of ones and m, n index the elements of J , ensuring that multiplication with q_{ij} results in a matrix filled with the corresponding penalty value. Next, we compute $\mathbf{M}_{k \times mn} = \mathbf{D}_{Q_k} + \mathbf{P}_{mn}$. The resulting \mathbf{M} is 33×33 , with 32 tooth classes (l) normalized to $(0,1)$. A background class (b) is added, with $b = 2$ to strongly penalize tooth-to-background misclassification, followed by final normalization for consistency. We integrate the geometrical and morphological prior of M within the loss function as follows:

$$L_{\text{GeoWDL}}(\hat{\mathbf{p}}, \mathbf{p}) = 1 - \frac{2 \sum_l \sum_i \mathbf{p}_{i,l} (1 - W^M(\hat{\mathbf{p}}_i, \mathbf{p}_i))}{2 \sum_l [\sum_i \mathbf{p}_{i,l} (1 - W^M(\hat{\mathbf{p}}_i, \mathbf{p}_i))] + \sum_i W^M(\hat{\mathbf{p}}_i, \mathbf{p}_i)}, \quad (2)$$

where $W^M(\hat{p}_i, p_i)$, given as: $W^M(\hat{\mathbf{p}}_i, \mathbf{p}_i) = \sum_{l=1}^L p_{i,l} \sum_{l'=1}^L M_{l,l'} \hat{p}_{i,l'}$ is the WD-weighted probability mass between predicted \hat{p}_i and ground truth (GT) p_i at voxel i . To address third molar class imbalance, the final segmentation loss L_{seg} combines L_{GeoWDL} with inverse class frequency weighted cross-entropy L_{WCE} .

Deep Watershed Instance Regression. To generate inputs for the deep watershed algorithm, our method optimizes two complementary tasks: energy basin regression and directional gradient estimation for boundary refinement. These tasks ensure accurate instance separation while encoding tooth boundaries and instance identity, enabling full tooth spatial understanding. We adapt the 2D deep watershed approach [2] to 3D tooth instance segmentation, modeling each tooth as an energy basin with a smooth energy gradient. First, we replace discrete energy-level classification with continuous energy map regression, which encodes spatial structure by assigning each voxel a distance to the nearest tooth boundary. This contrasts with the original approach and [9,29], which classify discrete offsets from the tooth centroid within a multi-task approach. Second, we refine boundary localization by estimating energy descent directions at each voxel, crucial for capturing rapid gradient changes in root apices. Unlike sequential pretraining [2], we train direction estimation as a parallel auxiliary task for efficiency. These adaptations aim to enable end-to-end optimization, improving instance awareness and refining segmentation precision, particularly in fine root structures. We compute watershed energy basins from GT using the Euclidean Distance Transform, which encodes distances to the instance boundary: $L_{EDT} = \frac{1}{N} \sum_{x=1}^X \sum_{y=1}^Y \sum_{z=1}^Z [I(x, y, z) - \hat{I}(x, y, z)]^2$. For energy direction, let $E(\mathbf{r})$, where $\mathbf{r} = (x, y, z)$, be the energy map defining a scalar field over a 3D voxel grid. The gradient $G(r)$ at each voxel r is computed by convolving the scalar field $E(r)$ with the 3D Sobel-Feldman operator K_d along each dimension d : $G(r) = \sum_{d \in \{x, y, z\}} K_d * E(r)$. The gradient magnitude $G(r)$ and the unit direction vector \mathbf{u}_v for each voxel v are given by $G(\mathbf{r}) = \|G(r)\|_2$ and $\mathbf{u}_v = \frac{G(r)}{G(r)}$, where $\|\cdot\|_2$ denotes the Euclidean norm. Maximum angular error $\theta_v = \pi$ occurs when \mathbf{u}_v misidentifies instance’s center, refining boundary localization. We optimize the 3-channel decoder’s D output using the mean squared error loss in the angular domain: $L_{dir} = \sum_{v \in P_l} \|\cos^{-1} \langle \mathbf{u}_{p_{GT}}, \mathbf{u}_{p_{pred}} \rangle\|^2$, where P_l and $l \in \{1, 2, \dots, 32\}$ is the set of all voxels belonging to the tooth semantic class. We mask non-tooth areas to reduce complexity and accelerate convergence. We clip \cos^{-1} to $\langle -1, 1 \rangle$, for numerical stability.

Deep Watershed Instance Classification via Majority Voting. Since instance segmentation assigns a single, consistent label to each tooth, we first apply the deep watershed algorithm to separate tooth instances and then classify them using majority voting based on voxel-wise predictions from the semantic segmentation branch. To this end, we extract instance seeds by empirically thresholding energy basins at half their depth ($\beta = 0.5$). Next, we binarize the multi-class segmentation to create a mask, restricting the watershed algorithm to tooth regions for improved computational efficiency. We then apply the watershed algorithm using predicted energy maps, seed points, and the segmentation mask, derived from an end-to-end optimized model, to separate 3D tooth instances. Finally,

Table 1. Quantitative results of GEPAR3D and state-of-the-art methods for general and tooth-specific segmentation. We report Detection Accuracy (DA) and F1 for instance detection and classification; DSC, RC, and HD for multi-class segmentation; and NSD_1 and RC_B for binary segmentation, with means and standard deviations (in brackets). Results are averaged across three external datasets unless stated explicitly. Methods are sorted by average DSC; best and second-best tooth-specific methods are highlighted in bold and underlined, respectively. **I**, **S**, and **IS** denote instance, semantic, and instance-based multi-class segmentation. † indicates p-value < 0.05.

Method	Type	DA (%)†	F1 (%)†	DSC (%)†				RC (%)†	HD (mm)↓	NSD_1 (%)†	RC_B (%)†
				In-house	Cui et al.	TF2	Average				
U-Net † [25]	S	95.6(4.6)	93.5(6.3)	87.8(3.4)	88.7(3.3)	88.3(3.1)	88.2(3.8)	85.5(5.0)	21.78(11.24)	88.2(5.1)	90.6(2.9)
Swin SMT † [23]	S	98.1(3.1)	96.8(4.0)	92.8(2.5)	92.9(2.5)	91.4(3.0)	92.3(2.8)	91.1(4.2)	2.93(1.89)	94.6(3.3)	92.9(3.2)
Swin UNETR † [30]	S	97.9(3.3)	96.6(4.6)	92.8(2.7)	92.6(2.3)	92.3(2.4)	92.6(2.7)	91.3(3.8)	3.41(2.57)	94.5(3.4)	93.3(2.7)
Swin UNETRv2 † [12]	S	98.1(3.3)	97.3(4.0)	92.7(3.2)	93.2(2.5)	93.4(1.8)	93.1(2.7)	91.6(4.3)	2.42(1.19)	95.5(3.4)	93.1(3.2)
ResUNet34 † [11]	S	98.4(3.5)	97.5(4.5)	93.5(2.1)	93.4(2.3)	93.0(2.7)	93.3(2.4)	90.6(3.9)	2.19(1.56)	96.0(2.9)	91.8(3.2)
VSmTrans † [20]	S	98.9(2.3)	97.7(3.6)	93.2(2.8)	93.5(1.7)	93.8(1.7)	93.5(2.3)	92.1(3.7)	9.06(7.91)	95.5(3.3)	94.1(2.5)
V-Net † [22]	S	98.9(2.5)	97.8(3.5)	93.7(1.7)	93.8(2.1)	93.2(2.3)	93.5(2.1)	92.4(3.6)	1.96(0.70)	95.9(2.9)	94.0(2.8)
Jang et al. † [13]	I	96.0(6.2)	-	83.5(1.6)	82.6(2.0)	82.5(1.3)	83.0(1.8)	75.6(6.1)	3.07(0.76)	79.3(3.0)	76.6(5.1)
MWTNet † [4]	I	92.6(8.4)	-	87.4(1.4)	84.3(2.0)	89.3(1.1)	87.4(2.5)	73.9(9.2)	2.29(0.59)	85.7(4.3)	76.3(6.2)
TSG-GCN † [21]	S	86.9(9.7)	83.0(11.4)	89.3(1.7)	91.0(3.4)	87.8(1.9)	89.2(3.0)	76.8(11.2)	2.47(0.76)	90.1(4.5)	86.3(4.9)
ToothSeg † [9]	IS	88.8(10.5)	86.2(7.5)	89.3(1.8)	93.6(0.8)	89.8(1.7)	90.4(2.6)	80.2(11.1)	2.84(1.67)	91.3(5.1)	81.0(8.0)
SGANet † [19]	S	92.9(9.6)	90.8(10.4)	92.2(1.6)	92.9(2.5)	91.9(1.8)	92.2(2.1)	83.8(9.4)	2.18(0.74)	94.3(3.2)	85.7(5.9)
GEPAR3D	IS	99.2(2.4)	98.0(3.7)	95.5(1.2)	95.1(0.8)	94.3(1.1)	95.0(1.4)	93.9(3.2)	1.44(0.70)	97.6(1.9)	95.2(2.1)

each instance is assigned the class with the most frequent voxel-wise prediction. For a given instance j , let C_i be the class of voxel i in segmentation S , and Iv_j its volume: $C_j = \operatorname{argmax}_c \sum_{i \in Iv_j} \delta(C_i, c)$, where $\delta(C_i, c)$ is 1 if $C_i = c$, otherwise 0. The instance is assigned the class with the highest count.

3 Experiments and results

Datasets and preprocessing. We train and validate our method on a publicly available dataset of 98 CBCT scans [7], reannotated into 32 classes following the Universal Numbering System [1]. We test on 46 CBCT scans from 4 medical centers, including two public datasets: Cui et al. [7] and Tooth Fairy 2 (TF2) [3,6] (file IDs in accompanying JSON), and 2 in-house sets from a retrospective study (IRB OKW-623/2022) at Polish centers A (11 scans, Carestream CS 9600) and B (9 scans, i-CAT 17-19). For reliable root evaluation, only scans with fully visible roots are included. All scans are resampled to 0.4 mm isotropic resolution, with Hounsfield Unit intensities clipped to $[0, 5000]$ and normalized to $[0, 1]$.

Implementation details. For training, we randomly crop 128^3 patches around GT-based ROIs. The model trains for 1000 epochs with AdamW, batch size of 2, and a cosine annealing scheduler. The loss function is defined as $L = A_1 L_{EDT} + A_2 L_{seg} + A_3 L_{dir}$, with empirically set weights $A_1 = 10$, $A_2 = 0.1$, $A_3 = 1e^{-6}$ for balance. The initial learning rate and weight decay are set to $1e^{-3}$ and $1e^{-4}$, respectively. During inference, a lightweight 3D U-Net (256³ patch) performs coarse binary segmentation to extract the ROI. Then we use sliding window inference (0.6 overlap, Gaussian weighting). The pipeline is implemented in PyTorch 1.13.1/MONAI 1.3.0 and runs on a single NVIDIA A100 GPU.

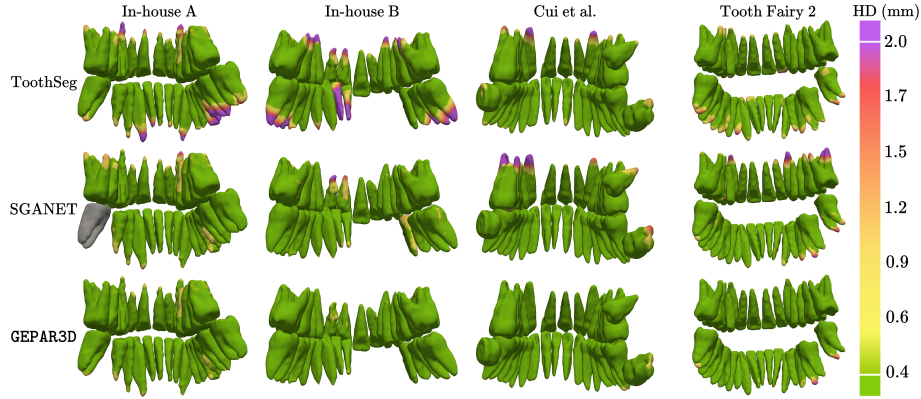


Fig. 3. Qualitative comparison of GEPAR3D with the two best-performing methods from quantitative results. Surface Hausdorff Distance heatmaps overlaid on GT labels (green = low, purple = high) highlight apex deviations. GEPAR3D shows superior root sensitivity versus tooth-specific baselines. Missing teeth are shown in gray.

Evaluation details. We evaluate on multi-class and binary metrics. Multi-class performance uses DSC, Precision (PR), Recall (RC), and Hausdorff Distance (HD). Binary evaluation uses Normalized Surface Dice within a 1-voxel GT boundary (NSD_1) [27] and Binary Recall (RC_B). While multi-class metrics assess overall performance, binary metrics focus on tooth tissue segmentation completeness, with RC_B highlighting false negatives and NSD_1 measuring boundary accuracy, including roots. Instance detection is evaluated via Detection Accuracy (DA) at a 50% Intersection over Union (IoU) threshold, with instances having $\text{IoU} > 0.5$ considered detected. Classification performance is measured using the F1 score. We benchmark GEPAR3D against general segmentation and tooth-specific state-of-the-art methods. General models follow GEPAR3D’s training setup (32-class labels, sliding window inference) and tooth-specific methods follow original protocols, with preprocessing and augmentations matched where possible. Statistical significance is determined via a paired t-test ($p < 0.05$).

Comparison with state-of-the-art methods. As shown in Table 1, GEPAR3D surpasses all competing methods in segmentation and instance detection. It achieves a DA of $99.2 \pm 2.4\%$ (+3.2% over Jang et al.) and an F1 score of $98.0 \pm 3.7\%$ (+7.2% over SGANet), confirming its superior ability to identify and classify tooth instances. This strong detection performance ensures segmentation metrics remain representative, even in challenging cases. GEPAR3D achieves the highest DSC on all external test sets, averaging $95.0 \pm 1.4\%$ (+2.8% over SGANet), alongside the best RC ($93.9 \pm 3.2\%$) and lowest HD (1.44 ± 0.50 mm), demonstrating robust generalization. NSD_1 of $97.6 \pm 1.9\%$ (+3.3% over SGANet) and RC_B of $95.2 \pm 2.1\%$ (+9.5% over TSG-GCN), highlight superior tooth tissue completeness. Qualitative results (Fig. 3) further validate this, revealing that competing methods often miss substantial root fragments, as shown by per-voxel

Table 2. Ablation study on network and loss components. The best-performing method is highlighted in bold, and the second-best is underlined. DA indicates detection accuracy, PR precision, RC recall and NSD normalized surface dice within 1 voxel boundary. **G** denotes Geometric Prior loss, **E** Energy map and **D** Direction map. $U_{0,1}$ denotes uniform distribution of random cost matrix. † indicates p-value < 0.05.

#	G	E	D	DSC (%)†	PR (%)†	RC (%)†	NSD_1 (%)†	DA(%) †	F1 (%)†
1†	-	-	-	93.27(2.40)	94.59(3.73)	90.62(3.85)	95.95(2.94)	98.4(3.5)	97.5(4.5)
2† ($U_{0,1}$)	-	-	-	93.49(2.35)	90.78(5.31)	92.76(3.53)	95.96(2.83)	99.2(2.0)	98.2(3.3)
3†	✓	-	-	94.55(1.49)	91.67(5.39)	92.86(2.35)	96.05(1.94)	99.3(1.9)	98.4(3.2)
4†	-	✓	-	94.58(1.28)	95.55(2.94)	91.36(3.67)	97.25(1.93)	99.2(2.2)	98.0(3.5)
5†	-	✓	✓	94.68(1.13)	95.52(2.86)	91.65(3.81)	97.41(1.83)	99.2(2.4)	98.0(3.5)
6†	✓	✓	-	94.96(1.13)	94.33(3.24)	93.12(3.33)	97.58(1.90)	99.1(2.5)	97.9(3.6)
7	✓	✓	✓	95.01(1.36)	94.95(3.45)	93.90(3.18)	97.63(1.94)	99.2(2.4)	98.0(3.7)

HD heatmaps, whereas GEPAR3D preserves more complete tooth anatomy.

Ablation Study. We evaluate the impact of various loss functions and network components in GEPAR3D (Table 2). As a baseline (#1), we use Dice+WCE loss. In (#2), we use the original GWDL with a penalty matrix randomly generated from a uniform distribution to test the loss function’s robustness to an uninformative prior (UP). Driven purely by error minimization via OT, shifts the PR–RC balance toward higher sensitivity, resulting in an RC increase of +2.14% while PR decreases by -3.81%. Despite the UP, DSC improves (+0.22%), confirming that loss regularization is not rigid and still enables the model to learn useful representations. Introducing the proposed geometric prior GeoWDL (G) in (#3) enhances DSC (+1.06%), improves NSD (+0.09%), and achieves the highest DA of 99.3%, demonstrating that structured guidance better aligns with tooth classification, though lowered PR remains. Adding energy map regression (E) in (#4) via the deep watershed method improves DSC (+1.31%) and boosts PR (+0.96%), thereby enhancing focus on tooth instances. In (#5) we incorporate an auxiliary energy descent direction task (D), yielding further gains in DSC (+0.10%), RC (+0.29%), and NSD (+0.16%), suggesting refined boundary localization. The introduction of GeoWDL in (#6) increases DSC (+0.28%) and noticeably improves RC (+1.47%). Finally, the proposed solution (#7), which jointly optimizes E and D under G guidance, not only raises PR (+0.36%) but, more importantly, significantly boosts RC (+3.28%) over (#1) and achieves the highest NSD of 97.63%. Overall, these results demonstrate that proposed components complement each other, enhancing sensitivity in challenging regions.

4 Conclusions

We present GEPAR3D, which combines geometric prior-assisted learning with deep watershed instance detection to improve tooth segmentation, particularly for fine root structures. Extensive experiments demonstrate its superiority over state-of-the-art methods, with enhanced segmentation supporting better orthodontic

planning and root resorption assessment. We ensure reproducibility by validating on public datasets, sharing code and implementation details. However, our study has limitations. Training was restricted to adult teeth, which may limit applicability to younger patients. Additionally, while geometric prior loss is crucial for encoding anatomical constraints, it can be overly sensitive when used alone, requiring careful tuning. In GEPAR3D, its integration with instance regression balances sensitivity and precision, mitigating this issue. While our method significantly improves root segmentation, further gains could be achieved with larger datasets and self-supervised training. Finally, resorption analysis requires comparing sequential scans to a reliable baseline segmentation. As no public CBCT datasets include resorbed annotations, we focused on validating apex segmentation accuracy, since under-segmentation would mask subsequent root shortening. To conclude, this work underscores the importance of root segmentation and aims to inspire future research.

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Disclosure of Interests. The authors have no competing interests to declare.

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