

Supplementary Information for: A fully automatic AI system for tooth and alveolar bone segmentation from cone-beam CT images

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Hierarchical data annotation method

We applied a hierarchical data annotation method to iteratively annotate all the CBCT scans in our dataset. Specifically, we first randomly selected 100 CBCT scans from the dataset, and 3 senior raters manually annotated the label including individual teeth and alveolar bones. This stage of fully manual annotation took about 2 months. Then, we used the 100 annotated CBCT scans to train our proposed system, and tested on the next 600 CBCT scans to produce initial segmentation results. Ten junior raters independently corrected the 600 labeled CBCT images to obtain preliminary results. Next, three senior raters checked the preliminary labeling results, and made the final decision. The second stage took about 5 days for correction. At last, the 700 annotated CBCT scans were used to train the AI system, which in return tested on the rest 4238 CBCT scans. Again, the final labeling results are corrected and checked by the ten junior raters and three senior raters, which took about 40 days to complete the process. The flow of the hierarchical data annotation method is shown in Supplementary Fig. 1.

Instead, to avoid inter-practitioner's variations in the data annotation process, all CBCT scans were independently labeled or checked by the 3 senior raters. When disagreements occurred among the 3 senior raters, they would have a discussion to make the final decision.

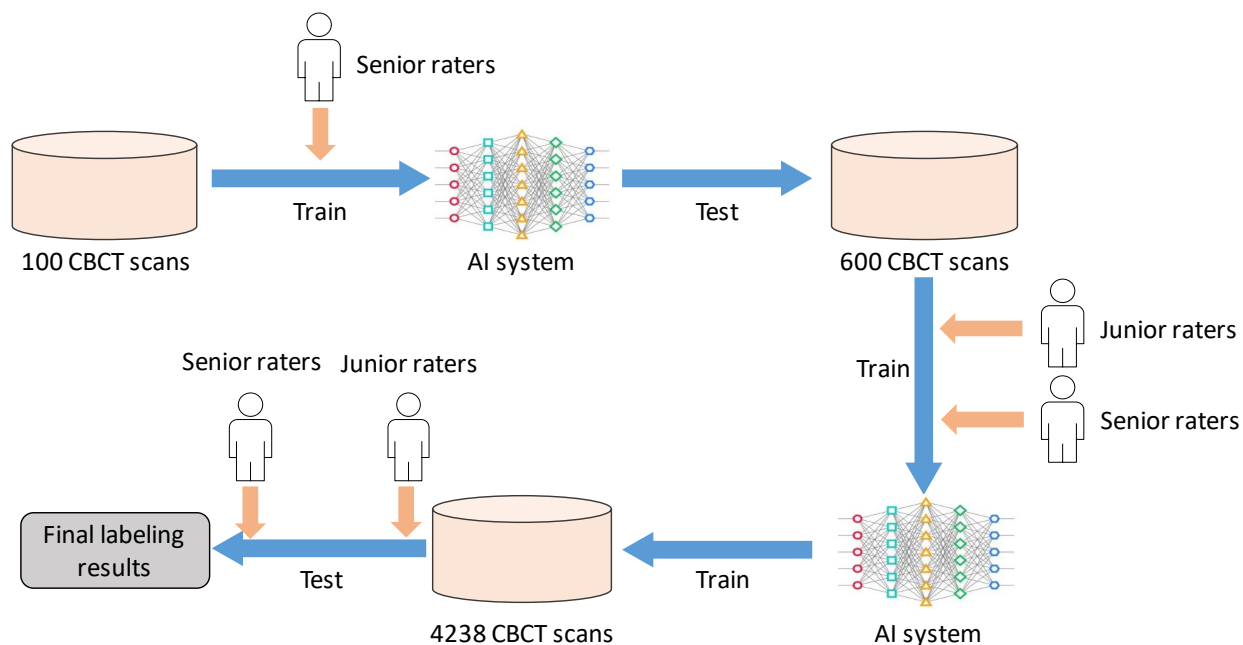
Dental abnormalities	Internal set		External set
	Training dataset (3172 CBCT scans)	Internal testing dataset (1359 CBCT scans)	External testing dataset (407 CBCT scans)
Missing teeth	946	345	137
Misalignment	2351	1085	314
Metal artifacts	435	199	96

Class	Model	Internal testing set			External testing set		
		Dice (%)	Sen (%)	ASD (mm)	Dice (%)	Sen (%)	ASD (mm)
Tooth	Ours (w/o S)	91.8	91.6	0.19	91.1	89.9	0.24
	Ours (w/o M)	92.7	91.9	0.23	91.4	90.3	0.29
	Ours	94.1	93.9	0.17	92.5	92.1	0.21
Bone	Ours (w/o H)	91.8	89.4	0.54	90.2	88.3	0.67
	Ours	94.5	93.8	0.33	93.8	93.5	0.40

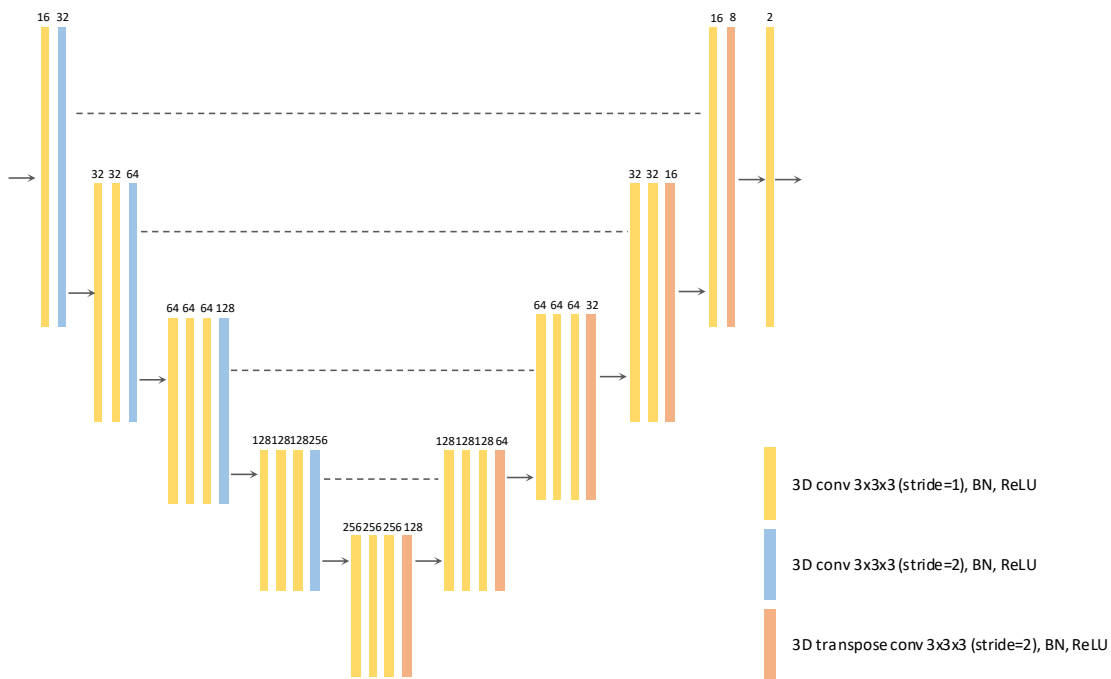
Ours (w/o S): our AI system without skeleton information in the inputs of the 2nd-stage tooth segmentation network.
Ours (w/o M): our AI system without the multi-task learning scheme, and only predicting the tooth mask of the 2nd-stage tooth segmentation network.
Ours (w/o H): our AI system without the harr filter enhancement for bone segmentation.

Methods	Training set (cases)	Dice (Tooth) (%)	Dice (bone) (%)
ToothNet* ¹	100	84.9	-
	100+	85.4	-
	3172	90.6	-
MWTNet* ²	100	83.7	-
	100+	85.6	-
	3172	90.4	-
CGDNet* ³	100	85.8	-
	100+	87.2	-
	3172	91.0	-
Ours	100	87.6	88.0
	100+	88.3	88.9
	3172	92.5	93.8

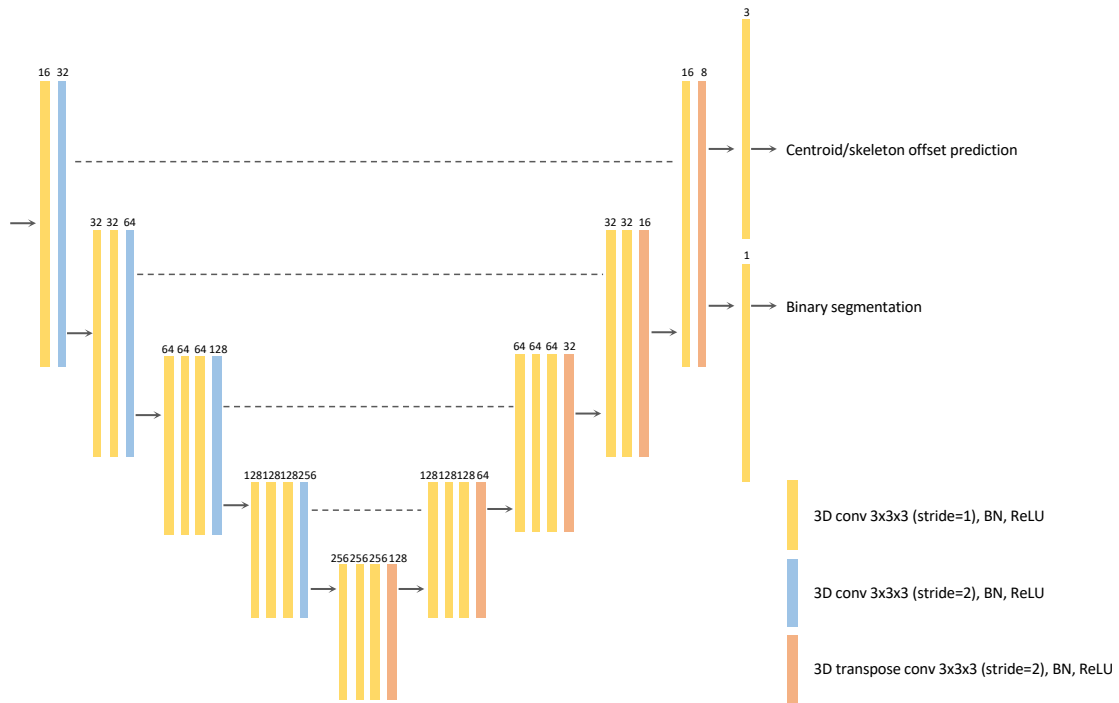
“*” means the method needs manual intervention (i.e., manually delineate foreground dental ROI). “+” means data augmentation techniques are used, including image flip, rotation, random deformation, and the learning based generative model.



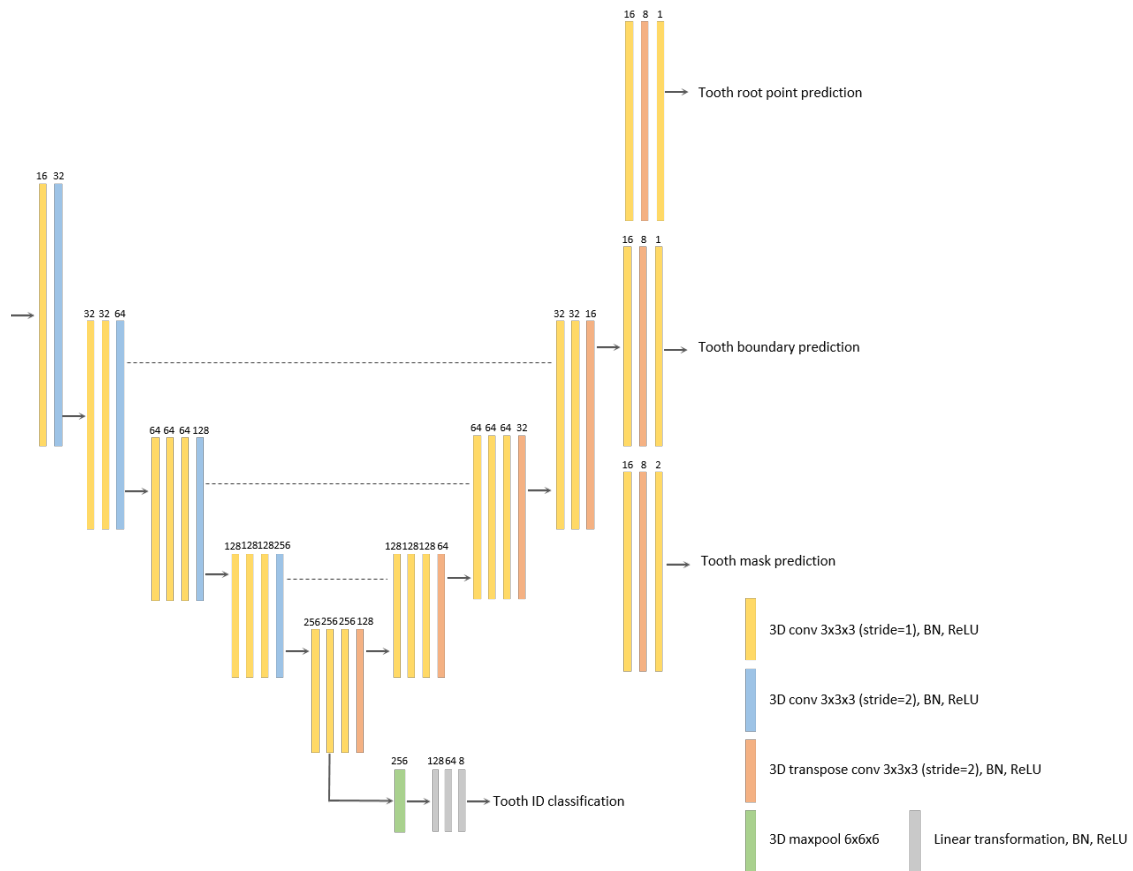
Supplementary Figure 1. The work flow of our hierarchical data annotation process.



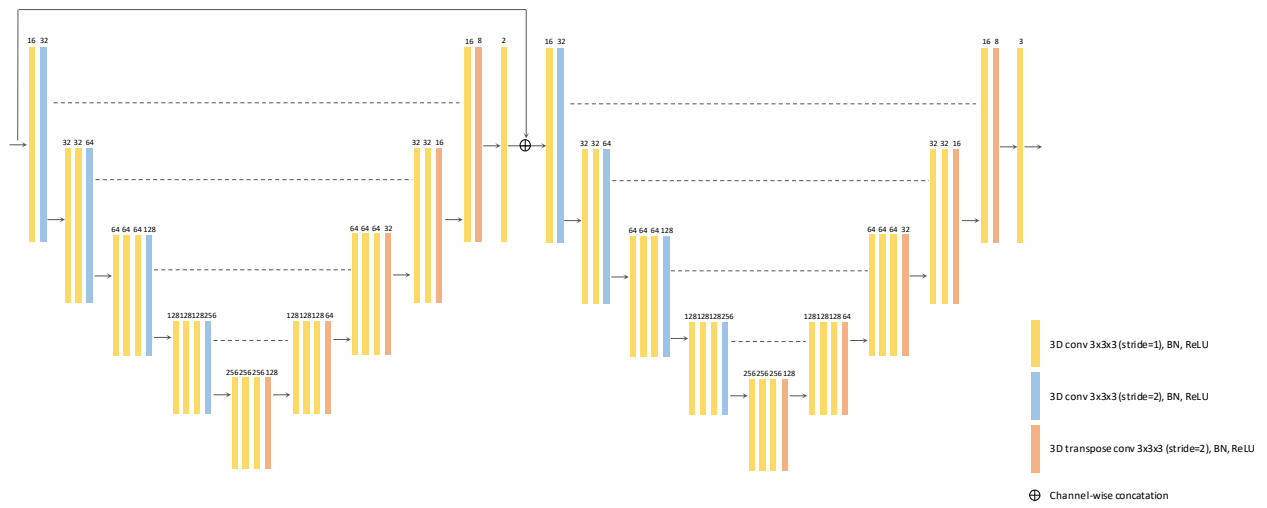
Supplementary Figure 2. The detailed network architecture of the ROI extraction network.



Supplementary Figure 3. The detailed network architecture of the tooth centroid or skeleton prediction network.



Supplementary Figure 4. The detailed network architecture of the multi-task tooth segmentation network.



Supplementary Figure 5. The detailed network architecture of the cascaded bone segmentation network.

References

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