

A Deep Learning Model to Predict Dose-Volume Histogram of Organs at Risk in Radiotherapy Treatment Plans

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INTRODUCTION

There are deep learning models^{1–8} including our previous studies^{2,8} can predict three-dimensional dose distributions, but none of them can directly predict clinically relevant DVHs for OARs. In this work, we would like to propose a deep learning method to directly predict achievable DVHs for OARs for automation of inverse planning. Based on what we known, this is for the first time to demonstrate a deep learning model to directly predict DVHs for OARs. The model was trained and evaluated with treatment plans of nasopharyngeal cancer cases. The deep learning architecture used is similar to our previous study⁸ but with different feature maps in each convolution layers. Compared with UNet, this architecture applies a residual network to perform down-sampling, each down-sampled part then being combined with deconvolution to perform up-sampling. Here, we firstly propose a concept of dose-area histogram as a bridge between a DVH and the deep learning network.

AIM

To develop a deep learning-based model to predict achievable dose-volume histograms (DVHs) of organs at risk (OARs) for automation of inverse planning.

METHOD

- 230 nasopharyngeal cancer patients (170 for training, 60 for testing) were included.
- The model was based on a connected residual deconvolution network (CResDevNet) and compared with UNet as a baseline.
- The input data comprised four channels for each OAR, and the output data is dose-area histogram (DAH) for that OAR:

$$DAH(D) = 1 - \frac{1}{A} \int_0^{D_{max}} \frac{dA(D)}{dD} dD$$

- The model was trained from scratch by correlating anatomical features with DAH of OARs, then accumulating these histograms to obtain DVH for each OAR.
- The DVHs and dose-volume indices (DVIs) for each OAR in the testing dataset were predicted to evaluate the accuracy of the models. Dice similarity coefficients for areas of the DVHs were also evaluated.

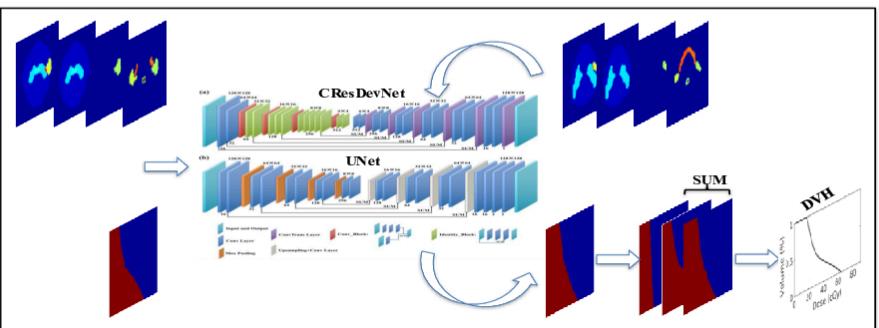


Fig. 1. Overview of the workflow for predicting the DVH for OAR.

RESULTS

- DVHs of 21 OARs in nasopharyngeal cancer were predicted using the models as shown in Fig. 2.
- For each patient, 63 DVIs for all OARs were calculated. Using the 60 patient treatment plans in the testing dataset, 79% and 73% of the DVIs predicted using the CResDevNet and UNet models, respectively, were within 5% of the clinical values. (Table 1)
- The median value of the DVIs' mean absolute errors was $3.2 \pm 2.5\%$ and $3.7 \pm 2.9\%$ for the CResDevNet and UNet models, respectively. (Table 2)
- The average dice similarity coefficient (DSC) for all OARs was 0.965 using the CResDevNet model and 0.958 using the UNet model. (Table 3)

Table 1. The proportion of DVIs having predicted values within 5% and 10% error.

	Within 5% error (%)		Within 10% error (%)		p-value
	CResDevNet	UNet	CResDevNet	UNet	
Brain Stem PRV	93	87	100	100	
Brain Stem	86	88	99	98	
Spinal Cord PRV	94	94	97	97	
Spinal Cord	96	94	99	99	
Parotid L	88	67	99	95	
Parotid R	82	77	97	97	
Thyroid Gland	81	73	96	94	
Lens L	98	98	98	98	
Lens R	98	100	100	100	
Optic Nerve L	58	52	89	78	
Optic Nerve R	53	52	86	77	
Optic Chiasm	52	40	78	73	
Larynx	79	78	95	95	
Trachea	87	70	97	95	
Mandible L	87	83	98	98	
Mandible R	80	76	95	91	
TMJ L	58	55	81	79	
TMJ R	68	64	87	84	
Temporal Lobe L	79	74	100	97	
Temporal Lobe R	79	73	97	97	
Pituitary	68	42	92	75	

Table 3. The DSC of DVHs predicted using CResDevNet and UNet models.

	DSC		p-value
	CResDevNet	UNet	
Brain Stem PRV	0.97±0.018	0.97±0.017	0.272
Brain Stem	0.963±0.025	0.964±0.028	0.818
Spinal Cord PRV	0.964±0.026	0.961±0.027	0.379
Spinal Cord	0.975±0.025	0.972±0.028	0.005
Parotid L	0.971±0.018	0.956±0.026	0.000
Parotid R	0.970±0.017	0.962±0.021	0.009
Thyroid Gland	0.952±0.011	0.961±0.020	0.000
Lens L	0.931±0.074	0.917±0.077	0.000
Lens R	0.926±0.050	0.919±0.041	0.225
Optic Nerve L	0.962±0.037	0.950±0.043	0.004
Optic Nerve R	0.959±0.027	0.950±0.031	0.075
Optic Chiasm	0.937±0.049	0.935±0.042	0.657
Larynx	0.971±0.018	0.966±0.027	0.033
Trachea	0.978±0.012	0.974±0.014	0.032
Mandible L	0.984±0.010	0.980±0.012	0.001
Mandible R	0.979±0.019	0.976±0.021	0.043
TMJ L	0.970±0.016	0.966±0.018	0.222
TMJ R	0.976±0.012	0.972±0.016	0.041
Temporal Lobe L	0.965±0.019	0.958±0.026	0.026
Temporal Lobe R	0.970±0.020	0.960±0.022	0.001
Pituitary	0.968±0.031	0.952±0.035	0.001

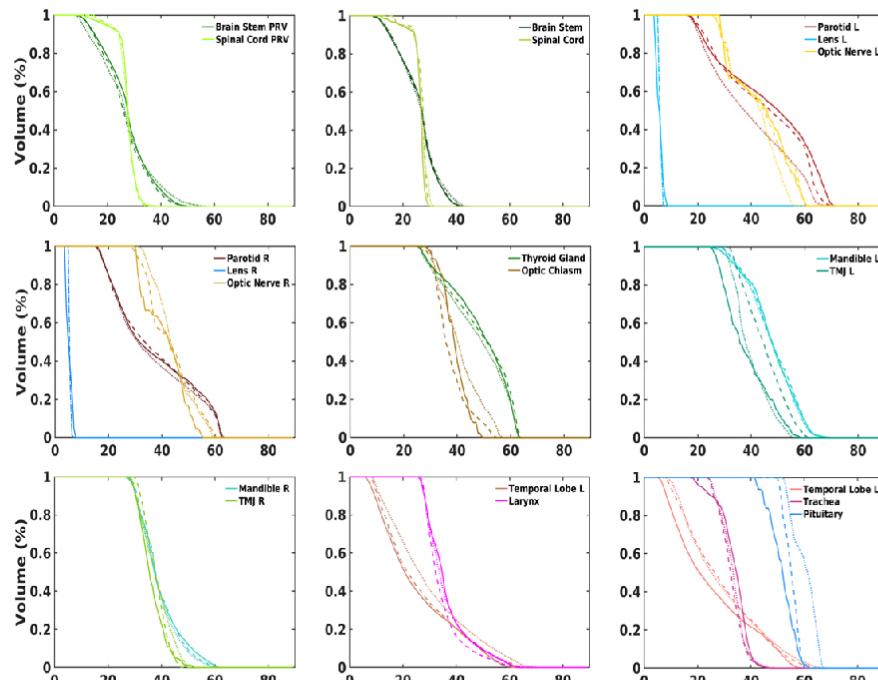


Fig. 2. Example of the "best" similarity between clinical (solid line), CResDevNet (dashed line), and UNet (dotted line) predictions of DVHs for each OAR.

Table 2. Mean absolute error (MAE) of DVIs predicted using CResDevNet and UNet models, together with their statistical comparison for each OAR.

	DVIs			p-value
	MAE (%) CResDevNet	MAE (%) UNet	p-value	
Larynx	V40 (%) 4.5±4.8	4.8±5.6	0.469	
D2%	2.9±2.5	3.4±2.7	0.173	
Dmean	2.5±1.8	2.5±2.0	0.965	
V40 (%) 2.9±3.4	4.8±5.0	0.001		
D2%	3.2±2.7	4.3±3.5	0.016	
Dmean	1.8±1.5	2.1±1.8	0.223	
Mandible L	V40 (%) 4.1±3.2	3.9±3.2	0.604	
V50 (%) 3.2±2.4	3.8±2.5	0.032		
D2%	1.9±1.7	2.7±2.1	0.003	
Dmean	1.1±1.1	1.4±1.2	0.022	
Mandible R	V40 (%) 4.4±3.7	5.1±4.6	0.069	
V50 (%) 4.7±4.6	5.3±5.2	0.089		
D2%	1.5±1.8	2.4±2.6	0.000	
Dmean	1.4±1.3	1.8±1.5	0.009	
TMJ L	V40 (%) 11.0±8.1	11.7±8.7	0.520	
V50 (%) 6.9±6.1	7.4±7.1	0.520		
D2%	3.4±2.6	3.3±3.0	0.836	
Dmean	3.2±2.1	3.4±2.5	0.499	
TMJ R	V40 (%) 9.0±7.8	9.2±7.5	0.843	
V50 (%) 5.1±5.1	5.5±5.3	0.541		
D2%	2.9±2.3	3.3±3.0	0.401	
Dmean	2.4±1.6	2.6±2.1	0.581	
Temporal Lobe L	V40 (%) 4.1±2.7	4.5±3.8	0.214	
V50 (%) 3.6±2.5	4.0±3.3	0.169		
D2%	3.3±2.5	3.7±2.9	0.255	
Dmean	2.0±1.9	2.6±2.3	0.005	
Temporal Lobe R	V40 (%) 3.2±2.4	3.8±2.5	0.038	
V50 (%) 2.6±2.0	3.9±2.9	0.000		
D2%	4.0±3.1	3.9±2.7	0.708	
Dmean	2.3±1.7	2.6±1.7	0.196	
Pituitary	D2% 4.4±3.8	7.1±4.9	0.000	
Dmean 4.5±5.0	7.1±5.4	0.001		

CONCLUSIONS

A deep learning model was built to directly predict achievable DVHs of OARs, taking a treatment site of nasopharyngeal cancer as an example. The model can automatically extract anatomical features of PTVs, OARs, and their geometric relationship. The predicted DVHs of OARs can be used for automation of inverse planning and quality assessment individual treatment plans.

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