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Inner Cascaded U²-Net: An Improvement to Plain Cascaded U-Net

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ABSTRACT

Deep neural networks are now widely used in the medical image segmentation field for their performance superiority and no need of manual feature extraction. U-Net has been the baseline model since the very beginning due to a symmetrical U-structure for better feature extraction and fusing and suitable for small datasets. To enhance the segmentation performance of U-Net, cascaded U-Net proposes to put two U-Nets successively to segment targets from coarse to fine. However, the plain cascaded U-Net faces the problem of too less between connections so the contextual information learned by the former U-Net cannot be fully used by the latter one. In this article, we devise novel Inner Cascaded U-Net and Inner Cascaded U²-Net as improvements to plain cascaded U-Net for medical image segmentation. The proposed Inner Cascaded U-Net adds inner nested connections between two U-Nets to share more contextual information. To further boost segmentation performance, we propose Inner Cascaded U²-Net, which applies residual U-block to capture more global contextual information from different scales. The proposed models can be trained from scratch in an end-to-end fashion and have been evaluated on Multimodal Brain Tumor Segmentation Challenge (BraTS) 2013 and ISBI Liver Tumor Segmentation Challenge (LiTS) dataset in comparison to related U-Net, cascaded U-Net, U-Net++, U²-Net and state-of-the-art methods. Our experiments demonstrate that our proposed Inner Cascaded U-Net and Inner Cascaded U²-Net achieve better segmentation performance in terms of dice similarity coefficient and hausdorff distance as well as get finer outline segmentation.

KEYWORDS

Deep neural networks; medical image segmentation; U-Net; cascaded; convolution block

1 Introduction

The first deep neural network applied in image segmentation field is Fully Convolutional Network (FCN) proposed by Long et al. in 2015 [1]. FCN tackles the challenge of semantic image segmentation by classifying each pixel as target or background and can accept input images of any size, whose variants get excellent segmentation performance on natural image segmentation field. However, the semantics of medical images are simpler compared to natural images and the structures are relatively fixed. The greatest difficulty in medical image segmentation field is the acquisition and labels of medical images, leading to a relatively small dataset. Thus, overfitting will be caused if the model



is too large and contains overmuch parameters. The most well-known deep learning baseline model in medical image segmentation field is U-Net proposed by Ronneberger et al. in 2015 [2]. The main differences between FCN and U-Net can be summarized as two aspects, one is a symmetrical encoderdecoder structure of U-Net which combines the low-resolution features in the encoder and the highresolution features in the decoder, the other is skip connections in U-Net to provide supplementary information in the up-sampling operations. And the model size of U-Net is relatively smaller. These features make U-Net been the first choice for medical image segmentation.

Since then, a mass of variants of U-Net have been proposed to boost its performance in various kinds of medical scenarios, including but not limited to improvements to skip pathways [3,4], more powerful and sophisticated U-blocks [5–9], cascaded U-structures [10–16], combining with attention mechanism [17–20] and so on.

Zhou et al. (UNet++) [3] proposed to redesign the skip connections using a sequence of conventional blocks to fill the gaps as opposed to straight connections in U-Net, which aims to decrease the semantic gap between the encoder and decoder feature maps. Huang et al. [4] further improved with full-scale skip connections to get performance gain with fewer parameters compared to UNet++, where each convolution block in the encoder is connected to the convolution block in the decoder of the same or lower level.

Zhang et al. [5] added residual connections in each convolution block, aiming to solve the problem of gradient vanishing and thus allowing to train deeper network. Ibtehaz et al. [6] propose MultiRes block to replace the two continuous convolution blocks. Inspired by Inception block [8], a sequence of three layers with 3×3 filters is applied to approximate 5×5 and 7×7 convolution operation, which is introduced to predict targets at different scales. Qin et al. [7] (U²-Net) developed a two-level nested U-structure for salient object detection by replacing conventional two convolutions in each level of U-Net with a residual U-block to extract intra-stage multi-scale features while remaining comparable memory and computation cost.

The most intuitive and simplest method to improve the segmentation performance is cascading several same or different U-Nets. The representation of cascaded U-Net architecture can be grossly divided into two groups, one is segmenting regions of interest firstly followed by target segmentation [10,11] and the other is rough pre-segmentation followed by explicit segmentation [12–16]. Christ et al. [10] first applied cascaded U-Net for automatic liver and lesion segmentation, in which the predicted liver ROIs segmented by the former U-Net are fed into the latter one as inputs to segment lesions only. Liu et al. [11] proposed Cascaded Atrous Dual-Attention UNet to leverage the interslice context information and emphasize the salient features of tumors. 3D AU-Net is first employed to get the coarse localization of the liver, whose prediction results are then concatenated with the input 3D feature map to preserve the z-axis information, followed by 2D ADAU-Net for precise tumor segmentation. Jiang et al. [12] proposed a two-stage cascaded 3D U-Net concentrating on the segmentation of substructures of brain tumors from rough to fine. Besides the coarse segmentation map from the first-stage U-Net, the raw images are fed together into the second U-Net too. Qin et al. [13] devised BASNet consisting of two U-Net-like networks adopted for salient object detection, with the prior U-Net learning to predict saliency map followed by residual refinement module which is also a U-Net refining the boundary. Li et al. [14] proposed a hybrid two-stage densely connected U-Net named H-DenseUNet for liver and tumor segmentation. The network includes a 2D DenseUNet for extracting intra-slice features due to memory limitation of 3D volume training and a 3D DenseUNet for extracting inter-slice features, which are fused eventually through a hybrid feature fusion layer. Liu et al. [15] devised a cascaded U-Net with residual mapping named CR-U-Net for liver segmentation,

in which morphological techniques, including opening and closing operation are used between the two U-Nets as an intermediate-processing module to refine the boundary. Hu et al. [16] proposed a multicascaded CNN to capture multi-scale features on brain MRI images with multi-modality. Specifically, three sub-networks are trained separately to segment the brain tumor from the axial, coronal, and sagittal views, respectively. The segmentation results are then fused to get the final segmentation masks.

1.1 Motivation

UNet++ [3] proposed in 2018 replaces the initial straight skip connections with a series of nested convolution blocks. Inspired by UNet++, we present Inner Cascaded U-Net, an improved cascaded network that places the encoder of the latter U-Net between the encoder and decoder rather than the end of the former U-Net. The inputs of each except the first and the last convolution block of the latter U-Net come from three aspects, which are the output from the previous layer of the same U-Net, the output from the former U-Net of the same level, and the corresponding up-sampled output of the lower convolution block of the former U-Net, respectively. By involving more between connections, the two U-Nets could share their learned features and the contextual information learned by the former U-Net can be incorporated into the latter one to enhance feature selecting and combination for better segmentation performance.

To further boost performance of Inner Cascaded U-Net, we propose Inner Cascaded U²-Net, which adopts residual U-block (RSU) proposed by Qin et al. [7] to replace the sequence of two plain convolution layers. A RSU block is a small U-Net like structure with residual connections, which can capture intra-stage multi-scale features while the residual connections allow to train deeper networks.

1.2 Contribution

There are many improvements to plain U-Net from various aspects, among which cascaded U-Net adopts the most intuitive method to cascade two successive U-Nets for the consideration of extracting features from coarse to fine. The main purpose of this article is to analyze the deficiency of the plain cascaded way and propose an improved inner cascaded method, which outperforms cascaded U-Net in segmentation accuracy and outline refinement. The contributions of the paper can be summarized as follows:

- We analyze different ways to improve the network architecture of the baseline deep learning model U-Net and the drawback of plain cascaded way.
- We propose a novel U-Net like architecture named Inner Cascaded U-Net, which organizes two U-Nets in an inner cascaded way and adds more between connections.
- We further propose Inner Cascaded U²-Net, which distinguishes from Inner Cascaded U-Net by residual U-block as an improvement to plain convolution block.

2 Proposed Methods

In this section, we first analyze the problem faced by plain cascaded U-Net. Then, we introduce Inner Cascaded U-Net, an improved combination method of two successive U-Nets. The inputs of the encoder and decoder convolution block of the latter U-Net are defined, respectively. Finally, we refer residual U-block and incorporate it into the network to get Inner Cascaded U²-Net.

2.1 The Problem of Plain Cascaded U-Net

The conventional cascaded U-Net combines a sequence of two U-Nets in an end-to-end way to predict feature maps from coarse to fine. The latter U-Net plays a role in improving outline segmentation of the former U-Net by inputting roughly segmented image masks. As shown in Fig. 1, the red arrow between the two U-Nets denotes the between connection where the input feature maps of the latter U-Net are generated by the former one. However, beyond that there are no other connections of the two U-Nets, which means the features that the former U-Net extracted cannot be properly shared with the latter one.



Figure 1: Illustration of cascaded U-Net architecture

2.2 Inner Cascaded U-Net

Enlightened by the nested dense skip pathways of UNet++, we design our Inner Cascaded U-Net to fuse the two successive U-Nets into a single U-Net like network, as shown in Fig. 2. Besides the down-sampled feature maps of the higher layer, the inputs of the encoder convolution block in the latter U-Net also involve the output of the convolution block of the same layer and the corresponding upsampled output of the lower layer in the former U-Net. Similarly, the inputs of the decoder convolution block comprise the output of the same layer in the former U-Net as well. UNet++ designs nested dense skip pathways in consideration of decreasing the semantic gap between the feature maps of the encoder and decoder. Here, we declare our purpose is to improve the cascade ways of two U-Nets by fusing more contextual information learned by the former U-Net into the latter one, instead of only taking the output of the former U-Net into consideration.

Formally, the inputs of the encoder convolution block which are represented as e in the latter U-Net can be formulated in Eq. (1):

$$e_{2}^{i} = \begin{cases} C\left(\left[e_{1}^{i}, U\left(e_{1}^{i+1}\right)\right]\right), & i = 1, \\ C\left(\left[e_{2}^{i-1}, e_{1}^{i}, U\left(e_{1}^{i+1}\right)\right]\right), & 1 < i < L, \\ C\left(\left[e_{2}^{i-1}, e_{1}^{i}\right]\right), & i = L, \end{cases}$$

$$(1)$$

where $C(\cdot)$ denotes the convolution operation followed by batch normalization and ReLU activation, $U(\cdot)$ is the up-sampling operation, and $[\cdot]$ represents the concatenation layer. L represents the number of layers of the U-Net. The superscript of e with $i \in [1, L]$ is the layer of U-Net, while the subscript of e with value of 1 or 2 denotes which U-Net e belongs to. As shown in Fig. 2, the inputs of the first convolution block in the latter U-Net come from only the former U-Net, and the last convolution block receives only inputs from down-sampled feature maps and output feature maps from the same level of the latter U-Net.



Figure 2: Illustration of our proposed inner cascaded U-Net architecture. The red and blue arrows denote added between connections from the former U-Net to the latter one

While the inputs of the decoder convolution block which are represented as d in the latter U-Net are described in Eq. (2):

$$d_2^i = C\left(\left[d_1^i, e_2^i, d_2^{i+1}\right]\right), \quad 1 \le i < L,$$
(2)

where all definitions are the same as encoder convolution block. Each encoder convolution block receives input feature maps from the corresponding encoder, the decoder of the former U-Net and the up-sampling feature maps from the lower decoder.

2.3 Inner Cascaded U²-Net

To further capture more global contextual information for better segmentation of small lesions in medical images, we propose Inner Cascaded U²-Net, which replaces the initial two successive convolution layers with Residual U-blocks (RSU) proposed in U²-Net [7]. The structure of RSU is shown in Fig. 3b, where H and W are the height and width of the input images, C_{in} and C_{out} are the input and output channels of feature maps, respectively. M denotes the channels of feature maps in the intermediate layers, which can be adjusted to control the augmented parameters. L represents the layers of the U-Net like network, while d is dilated convolution [21].

Compared to the plain convolution block shown in Fig. 3a, RSU replaces the second convolution layer with a U-Net like network whose output is then added to the output of the first convolution block, aiming to extract multi-stage features at small memory and computation cost while keeping a consistent output dimension with plain convolution block. Deeper levels with larger L lead to larger receptive field and thus more multi-stage contextual information can be extracted during progressive down-sampling and up-sampling process. However, larger L also means more encoder and decoder layers, leading to more memory and computation cost. L is set to 7 as recommend in [7].



Figure 3: Comparisons between the plain convolution block and RSU. (a) Origin plain convolution block in U-Net. (b) Residual U-block RSU

3 Experimental Results and Analysis

We conduct experiments and validate our methods on two public medical imaging datasets including brain MRI and liver CT. In this section, we first introduce the two datasets and preprocessing operations to each dataset, respectively. Then we present two evaluation metrics which are widely used in medical image segmentation field and the experiments environment in which we conduct our experiments. Finally, we give the performance analysis based on the quantitative and qualitative results.

3.1 Datasets and Preprocessing

The dataset for brain segmentation in MRI images is the Multimodal Brain Tumor Segmentation Challenge (BraTS) 2013 dataset. The dataset contains four modal images for each patient, which are T1, T1c, T2, Flair, respectively. It is composed of real and synthetic images and we only use the real images, including 20 high-grade gliomas (HG) patients and 10 low-grade gliomas (LG) patients. There are 4 different types of brain tumor in the dataset, which are all regarded as tumor positive for simplicity. The four modal images of the same size 240×240 are first resized to 256×256 using bilinear interpolation and then concatenated to one single image with four channels to enhance the segmentation performance. Considering the dataset is relatively small, it is randomly divided into training, validation and testing set at 60%, 20% and 20%, resulting in 3361, 1110, 1014 images, respectively.

The dataset for liver segmentation in CT images is the ISBI Liver Tumor Segmentation Challenge (LiTS) 2017 dataset. The dataset contains 131 patients so we use more data for training in view of the abundant images in the dataset, of which the first 100, the next 15 and the last 15 except the last

patients are used for training, validation and testing set, adding up to a total number of 13144, 3308 and 2518 images, respectively. The tumor areas are labelled as the liver for binary segmentation. And the CT values of the images in the dataset are preprocessed to liver window settings [-200, 250] to reduce distractions from other organs.

All images are normalized finally to set CT values between [0, 1] before inputting them to the network to accelerate the training process. The overall experiment pipeline on the BraTS 2013 and LiTS dataset is shown in Fig. 4, from which we can see the proposed architecture is trained in an end-to-end fashion with preprocessing steps as few as possible.



Figure 4: Overview of the proposed pipeline of brain tumor segmentation on the BraTS 2013 dataset and liver segmentation on the LiTS dataset. The operations within the dotted line denote the preprocessing steps

3.2 Evaluation Metrics

Dice Similarity Coefficient (DSC) and Hausdorff Distance (HD) [22] are used to evaluate the segmentation performance of our proposed method. DSC is widely used in medical image segmentation to measure the overlap degree between the predicted feature maps and ground truth, which is defined in Eq. (3):

$$DSC = \frac{2|P \cap G|}{|P| + |G|},\tag{3}$$

where P represents the predicted feature maps, G is the ground truth labels, $|\cdot|$ denotes the area of the region.

HD is a measurement to the similarity degree of two groups of point sets. And here are the predicted feature maps and ground truth labels. HD is conventionally multiplied by 95% (HD95) to

eliminate the influence of outliers. HD95 is defined in Eq. (4):

$$HD95 = \max\left(\max_{p \in P} \min_{g \in G} d(p,g), \max_{g \in G} \min_{p \in P} d(p,g)\right) * 0.95,$$
(4)

where *P* and *G* are the predicted feature maps and the ground truth labels, respectively, d(p, g) denotes the Euclidean distance.

3.3 Implementation Details

The implementation of our proposed model is based on Python 3.6 and Keras frame with Tensorflow backend. All experiments are conducted on a single machine with an Intel Core i9-10900X 3.70 GHz (64 GB RAM) and an NVIDIA RTX 2080Ti GPU (11 GB memory). In particular, the initial learning rate is set to 1e-4 and Adam optimizer is adopted to update the weights of the network, with learning rate decay of 0.1 and patience of 10 in LiTS or 4 in BraTS 2013 monitored on the loss of validation set. In the training process of LiTS dataset, the maximum training epochs is set to 20 and a mini batch size of 2 is applied due to limit of GPU memory, while the models are trained for 50 epochs using batch size of 8 for the training of BraTS 2013 dataset. All models in our experiments are trained from scratch with no data augmentation before training or post-processing after training.

Focal Tversky Loss (FTL) [23] rather than Dice loss is implemented in our experiments for the consideration of achieving a better tradeoff between precision and recall. FTL is defined in Eq. (5):

$$FTL = \left(1 - \frac{\sum_{i} p_{i}g_{i} + \varepsilon}{\sum_{i} p_{i}g_{i} + \alpha \sum_{i} p_{i}\left(1 - g_{i}\right) + \beta \sum_{i} g_{i}\left(1 - p_{i}\right) + \varepsilon}\right)^{\frac{1}{\gamma}},\tag{5}$$

where larger γ concentrates more on misclassified predictions and is set to its initial value 4/3. p_i denotes the predicted value between 0 and 1 of the *i*-th pixel. g_i denotes the true value of the *i*-th pixel, which is 0 (background) or 1 (brain tumor for BraTS 2013 or liver for LiTS dataset) in our experiments. α and β are hyperparameters used to balance precision and recall, which are set to 0.4 and 0.6, respectively. ε is implemented to prevent division by zero, here is set to 1e-6.

3.4 Quantitative and Qualitative Results

Table 1 compares the segmentation performance on BraTS 2013 and LiTS of our proposed Inner Cascaded U-Net and Inner Cascaded U²-Net with related 4 existing methods whose main ideas are referenced. As seen, cascaded U-Net outperforms U-Net on both datasets, which is the advantage of adding another U-Net with the same structure to refine the segmentation mask predicted by the single U-Net. While our proposed Inner Cascaded U-Net further improves the segmentation performance to cascaded U-Net by 0.5% and 0.2% respectively in DSC. Inner cascaded method distinguishes from plain cascaded U-Net by replacing the coarse-to-fine improved way with a single-stage method in which the latter U-Net plays a more important role in predicting where more contextual information learned by the former U-Net is incorporated into the latter U-Net to boost the overall segmentation performance. It is worth mentioning that Inner Cascaded U-Net gets the smallest value of HD95 on both datasets, which means the predicted results of Inner Cascaded U-Net is best-matched to the ground truth and shows the effectiveness of inner cascaded method. Our proposed Inner Cascaded U²-Net achieves the best performance in case of DSC on both datasets, which outperforms cascaded U-Net, U²-Net, Inner Cascaded U-Net by 0.6%, 0.5%, 0.1% respectively on BraTS 2013 and 0.9%, 1.1%, 0.7% respectively on LiTS. Standard convolution block consists of two successive convolution layers with 3 * 3 kernel, which is equal to a single convolution layer with 5 * 5 kernel [9] and thus is inadequate to extract multi-scale features especially the features of small lesions. While RSU in Inner Cascaded U^2 -Net is able to capture multi-scale features and incorporate more global contextual information through a small U-Net like network for better segmentation of small lesions, which further boosts the performance on medical image segmentation.

Method	BraTS 2013		LiTS		
	DSC	HD95	DSC	HD95	
U-Net	0.9396	2.39	0.9350	148.09	
Cascaded U-Net	0.9414	2.25	0.9433	66.12	
U-Net++	0.9340	2.68	0.9469	89.50	
U ² -Net	0.9429	2.02	0.9407	79.13	
Inner Cascaded U-Net (Ours)	0.9468	1.94	0.9452	46.39	
Inner Cascaded U ² -Net (Ours)	0.9474	2.07	0.9520	59.87	

 Table 1: Comparison of the proposed inner cascaded U-Net and inner cascaded U²-Net with other related methods on the BraTS 2013 and LiTS dataset in terms of DSC and HD95

The proposed Inner Cascaded U-Net and Inner Cascaded U²-Net are benchmarked against existing state-of-the-art methods on the BraTS 2013 and LiTS dataset in terms of DSC, as shown in Table 2. The results of BraTS 2013 dataset show that two cascaded methods including Cascaded DCNN and MCCNN achieve DSC of 0.8 and 0.89, respectively, while our proposed Inner Cascaded U²-Net outperforms the two methods a lot with DSC of 0.9474. The comparison results obtained from the LiTS dataset demonstrate our proposed Inner Cascaded U²-Net achieve the second highest DSC within the two-dimensional methods, among which CR-U-Net and Cascaded modified U-Net are designed in a cascaded method, which shows the effectiveness of Inner Cascaded U²-Net. Furthermore, two cascaded methods fusing 2D and 3D networks namely 2.5D methods are also presented in the table, which are H-DenseUNet and 3D AUNet + 2D ADAU-Net, respectively. The results demonstrate the 2.5D and 3D methods can lead to an enormous performance gain, which is due to the inherent defects in 2D models that they are incompetent to capture the spatial context information along z-axis, leading to segmentation performance degradation. Incorporating the inner cascaded method into three-dimensional models has not yet been implemented due to limitation by computing resources, which can be further researched.

Table 2: Comparison of the proposed inner cascaded U-Net and inner cascaded U ² -Net with the state
of-the-art methods on the BraTS 2013 and LiTS dataset in terms of DSC

Dataset	Method	Year	Dimension	DSC	
BraTS 2013	Cascaded DCNN [24] MCCNN [16] 3D dense connectivity network [25]	2017 2019 2020	2D 2D 3D	0.8 0.89 0.87	

(Continued)

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Dataset	Method	Year	Dimension	DSC
	Di-phase midway convolution and	2020	2D	0.91
	deconvolution network [26]			
	LSTM [27]	2020	2D	0.95
	Two-phase method [28]	2022	2D	0.9525
	Inner cascaded U-Net (Ours)	2022	2D	0.9468
	Inner cascaded U ² -Net (Ours)	2022	2D	0.9474
LiTS	ACM with FCN [29]	2019	2D	0.943
	CR-U-Net with dice loss [15]	2019	2D	0.9542
	HDA-ResUNet [30]	2021	2D	0.944
	Cascaded modified U-Net [31]	2021	2D	0.95
	H-DenseUNet [14]	2018	2.5D	0.961
	3D AUNet + 2D ADAU-Net [11]	2021	2.5D	0.9723
	Hybrid 3D residual network [32]	2020	3D	0.971
	Inner cascaded U-Net (Ours)	2022	2D	0.9452
	Inner cascaded U ² -Net (Ours)	2022	2D	0.9520



Figure 5: Qualitative comparison of proposed Inner Cascaded U-Net and Inner cascaded U²-Net with U-Net, cascaded U-Net, U-Net++ and U²-Net. The first two cases are visualization of segmentation masks from BraTS 2013 dataset, while the last two cases are from the LiTS dataset

Fig. 5 shows a qualitative comparison between our proposed models and other related 4 models. As we can see, our proposed Inner Cascaded U-Net and Inner Cascaded U²-Net can get more precise segmentation masks and finer outlines compared to other related models. Inner Cascaded U²-Net has

a better handle on the segmentation of different types of objects at different scales, which is attributed to the ability of RSU to capture multi-scale features. The second case from BraTS 2013 demonstrates promising segmentation performance of Inner Cascaded U²-Net, with other models all omitting the outline details of small objects. But, our Inner Cascaded U²-Net can still capture the main features and have a finer outline segmentation. Here the global contextual information extracted by RSU is utilized for boundary refinement. The last case from LiTS dataset shows less false positive predictions in UNet++ and our proposed methods compared to other three models, indicating the significance of shared contextual information. The single between connection in cascaded U-Net is insufficient to share the learned contextual information from the former U-Net to the latter one, leading to insensitivity to false positive predictions, which is improved by inner cascaded method where more contextual information is incorporated in each corresponding layer.

4 Conclusion

In this paper, we propose Inner Cascaded U-Net and Inner Cascaded U²-Net. Inner Cascaded U-Net refines the plain cascaded U-Net which connects in an end-to-end way by adding inner nested connections between the corresponding encoders and decoders of the two U-Nets, thus fusing more contextual information learned by the former U-Net into the latter one. Inner Cascaded U²-Net further enhances the segmentation performance by replacing the plain convolution block with residual U-block to capture more global contextual information for the sake of better segmentation of small regions. Experiments on the BraTS 2013 and LiTS datasets demonstrate that both models achieve excellent segmentation performance in terms of DSC, HD95 as well as refined segmentation masks.

Compared to the plain cascaded U-Net and U^2 -Net, however, the number of increased parameters of our proposed models are relatively large, leading to more computation and memory cost. Our future work will aim at exploring optimization strategy to decrease the model size and thus save the training as well as prediction time.

Availability of Data and Materials: The two datasets used to support our research are both publicly available. The BraTS 2013 dataset is available at https://www.smir.ch/BRATS/Start2013#!#download, accessed on 13 November 2021. And the LiTS dataset is available at https://competitions.codalab. org/competitions/17094#participate, accessed on 13 November 2021. All models mentioned in the experiments and code generated during the study by the corresponding author are available at https://github.com/FreedomXL/Inner-Cascaded-U-2-Net.

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Conflicts of Interest: The authors declare that they have no conflicts of interest to report regarding the present study.

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