

# Laplacian Saliency-Gated Feature Pyramid Network for Accurate Liver Vessel Segmentation

Zhan Gao, Qiuhaio Zong, Yiqi Wang<sup>1</sup>, Yan Yan, Yuqing Wang<sup>2</sup>, Ning Zhu, Jin Zhang<sup>3</sup>, Yunfu Wang, and Liang Zhao<sup>4</sup>

**Abstract**—Liver vessels generated from computed tomography are usually pretty small, which poses major challenges for satisfactory vessel segmentation, including 1) the scarcity of high-quality and large-volume vessel masks, 2) the difficulty in capturing vessel-specific features, and 3) the heavily imbalanced distribution of vessels and liver tissues. To advance, a sophisticated model and an elaborated dataset have been built. The model has a newly conceived Laplacian saliency filter that highlights vessel-like regions and suppresses other liver regions to shape the vessel-specific feature learning and to balance vessels against others. It is further coupled with a pyramid deep learning architecture to capture different levels of features, thus improving the feature formulation. Experiments show that this model markedly outperforms the state-of-the-art approaches, achieving a relative improvement of Dice score by at least 1.63% compared to the existing best model on available datasets. More promisingly, the averaged Dice score produced by the existing models on the newly constructed dataset is as high as  $0.734 \pm 0.070$ , which is at least 18.3% higher than that obtained from the existing best dataset under the same settings. These observations suggest that the proposed Laplacian saliency, together with the elaborated dataset, can be helpful for liver vessel segmentation.

**Index Terms**—Vessels, segmentation, computed tomography, neural network, pattern recognition and classification.

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Zhan Gao, Qiuhaio Zong, Yiqi Wang, Yan Yan, and Jin Zhang are with the Taihe Hospital, Hubei University of Medicine, Shiyan 442000, China (e-mail: gaozhan@taihehospital.com; zongqiuhaio@taihehospital.com; wangyiqi@taihehospital.com; yanyan@taihehospital.com; zhjinwhu@taihehospital.com).

Yuqing Wang and Ning Zhu are with the School of Biomedical Engineering, Hubei University of Medicine, Shiyan 442000, China (e-mail: 3347460663@qq.com; 1664891351@qq.com).

Yunfu Wang is with the Taihe Hospital/Hubei Key Laboratory of Embryonic Stem Cell Research, Hubei University of Medicine, Shiyan 442000, China (e-mail: yfwang@taihehospital.com).

Liang Zhao is with the Taihe Hospital/Hubei Key Laboratory of Embryonic Stem Cell Research, Hubei University of Medicine, Shiyan 442000, China, and also with the School of Computing and Electronic Informatics, Guangxi University, Nanning 530004, China (e-mail: S080011@e.ntu.edu.sg).

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## I. INTRODUCTION

LIVER vessel segmentation of medical images generated from computed tomography (CT) provides rich information for disease diagnosis, radiotherapy positioning and surgical planning [1]. Therefore, various computational models have been developed for this task, including traditional machine learning-based and deep learning-based.

The traditional methods mainly use image filters (e.g., Frangi [2], Hessian [3], Laplacian of Gaussian [4]), clustering algorithms [5], adaptive thresholding [6], active contour models [7], [8] to pinpoint out liver vessels. While the deep learning models are more prevalent and effective due to the revolutionary design of network architectures as well as the powerful devices. The fruitful innovations in this category are mainly two-fold: new loss functions and novel architectures. The former includes active contour loss [9], boundary loss [10], Dice loss [11], Tversky loss [12], etc.; while the latter attracts even more intensive efforts, e.g., the U-Net based models [13], [14], attention-based model [15], multi-path convolutional network [16], etc.

Although dozens of methods have been proposed from every perspective, the performance is still unsatisfactory. In fact, liver vessel segmentation has never been a trivial task because vessels usually have tiny size, low contrast and heavy noise, particularly the small size, which leads to the heavily skewed distribution of vessels versus other liver tissues. Taking Fig. 1, all the ratios between vessel pixels and liver pixels contained in the raw images are less than 0.1, some even less than 0.05, presenting great challenges for shaping vessel-specific feature learning. Another consequence of the smallness is the difficulty of formulating contextual features from vessel areas. Taking Fig. 1 again, many vessels only occupy dozens of, even less than ten pixels, resulting in very limited information to be retrieved. What's more, it is a painstaking task to delineate vessels from livers due to the tiny regions they occupy, hence the scarcity of large-volume and high-quality data. These challenges collectively lead to the poor performance of existing models.

To overcome these shortcomings, we propose a novel Laplacian-based saliency generator that is efficient and effective in highlighting vessel-like areas and suppressing others. For instance, by applying the filter to the liver images shown in Fig. 1, the averaged vessel-to-liver weighted pixel-level ratio is increased to 2.73 times higher than the original one. More

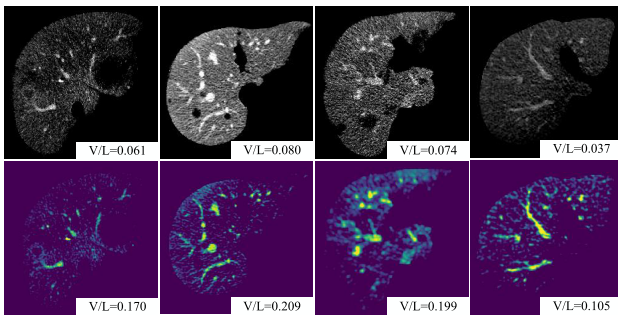


Fig. 1. Effectiveness of the proposed Laplacian salience filter. The first row contains the original images and the second is the Laplacian salience filtered ones. The value at the bottom right corner is the ratio between the weighted number of vessels and other liver tissues, where the weights are equal before filtering and are determined automatically by Laplacian salience after filtering.

importantly, the filter is insensitive to the phase of images, which can also be observed in Fig. 1. By integrating the filter into a pyramid deep learning architecture, the features can be more effectively tailored to the vessel areas, and the skewed distribution can be automatically balanced as well. In addition, the pyramid network is able to capture different levels of features and fuse them together for better vessel segmentation. The Laplacian salience automatically shapes the model focusing on vessel-like regions by attaching higher weights to these pixels and lower weights to others, thus paying more attention to vessel-like areas. For multi-scale feature fusion, it can adapt to various vessel sizes and compensate for weak features learned from small vessels by large vessels, so that the learned features can be applied to different situations.

Besides the sophisticated model, another big issue that hinders the progress of liver vessel segmentation is the small and imperfect dataset, which is also caused by the tiny size of vessels. To construct a qualified large dataset, great endeavors have been made by our team. In particular, 532 volumes of contrast-enhanced abdominal computed tomography images were collected, and the vessels were carefully delineated by three radiologists elaborately.

The big jump in Dice scores generated by our model, as well as from our data, demonstrates their goodness. The source codes and data are available at <https://github.com/lzhLab/LiVS>.

## II. RELATED WORKS

### A. Saliency Map

A saliency map highlights the focus of the human visual system, which has been demonstrated to be effective in dealing with inhomogeneous distributions as well as intra-class variations in visual tasks [17]. There have two typical branches related to saliency, namely saliency prediction and saliency application. Saliency prediction has been extensively studied from different perspectives, such as data-driven methods [18], [19], generative adversarial network-based models [20], [21], et al. Regarding saliency application, it is mostly used to facilitate downstream data analysis, where the saliency itself is mainly obtained by attention-based models and traditional machine learning-based methods. For instance, the

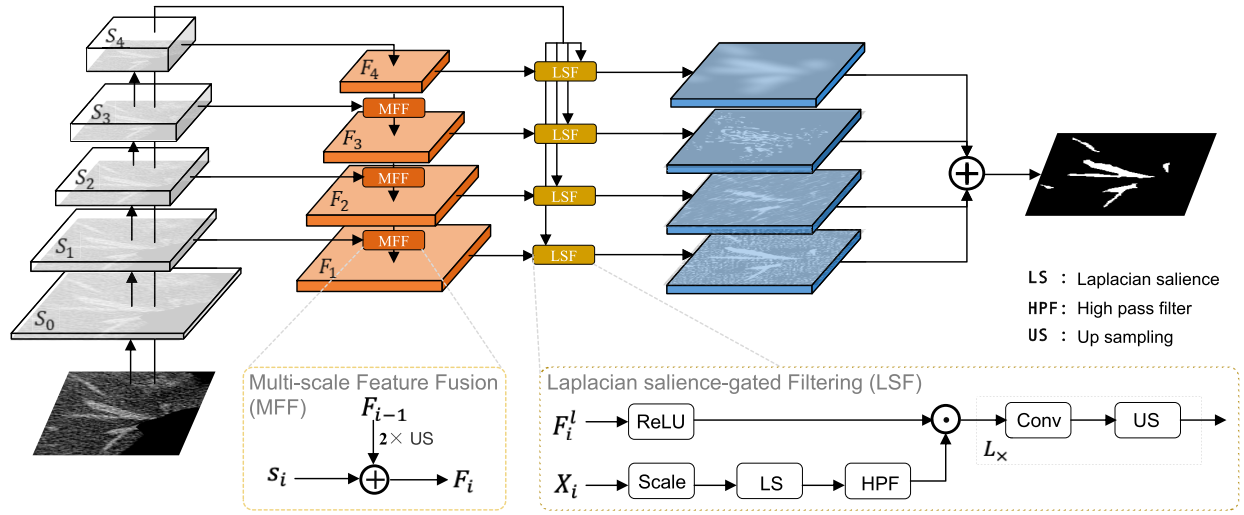
self-attention model [22] as well as its spatial variations [23], [24] and channel variations [25], [26] have been proposed to capture the saliency of objects. Another work, the attention gate [27], has also been proposed to suppress irrelevant regions and highlight salient features. This study is similar to ours in using saliency but different in generating saliency. Although these implicit model-driven saliency generators have been proven to be effective in real-world tasks, they require a large amount of high-quality training data. This restriction can be overcome by traditional approaches, such as Conditional Random Field-based [28], Bayesian inference-based [29], and clustering-based [30], but at the cost of accuracy.

### B. Loss Weighting

Loss weighting is dissimilar to saliency. It assigns different weights to different classes, samples, and even pixels, while saliency varies in weight across different regions, which may be composed of multiple classes. Loss weighting for image analysis can be categorized into three clusters typically, i.e., class-wise [31], [32], sample-wise [11], [12], [33], and pixel-wise [9], [10]. The class-wise weighting assigns different weights to different classes to balance the uneven distributions; the sample-wise weighting assigns different weights to different samples, even if they are from the same class, so that more attention could be paid to hard samples; and the pixel-wise weighting is the most fine-grained strategy, which can also apply different weights to different classes and samples. Among them, pixel-wise weighting is more popular. For instance, Focal loss [34] is a common weighting approach for different pixels based on the predicted probability, while contour loss [9] and boundary loss [10] have been adopted to focus on ambiguous regions across different classes.

### C. Tiny Object Segmentation

The cross-section of a vessel is a tiny blob since the branches of vessels are thin tube-like structures. Therefore, vessel localization in liver is similar to tiny object segmentation. Several intrinsic difficulties hinder the accurate segmentation of tiny objects, including information loss, noisy feature representation, insufficient samples and sensitive to disturbance [35]. To overcome these limitations, extensive efforts have been made from various perspectives [36], such as data augmentation [37], soft label assignment [38], scale-specific segmentation [39], feature reassembly [40], attention-based segmentation [41], similarity-aware learning [42], super-resolution-based segmentation [43], context-aware modeling [44], focus-aware segmentation [16], etc. The strategies presented here are not used separately, but in most cases, they are combined together to achieve better performance. For instance, Liu et al. [45] propose a dual-branch network with dual-sampling modulated Dice loss for hard exudate segmentation from color fundus images, in which each branch is an exudate size-specific feature learner so that more attention can be paid to small exudates. It also oversamples small exudates that are difficult to segment. In another study, Meng et al. [46] combine multi-scale feature fusion and the



**Fig. 2.** Architecture of the Laplacian salience-gated feature pyramid network. It is mainly composed of two parts, including pyramid feature fusion and the Laplacian salience-gated feature filtering. The pyramid network is for multi-scale feature extraction, while the Laplacian salience is to shape the vessel-specific feature learning. Note that the Laplacian salience is applied to different scales of the input image other than the corresponding feature maps, and the final feature map is the integration of the feature maps obtained from different scales with size matched.

Transformer for fine-grained segmentation of liver tumors and vessels.

Note that object segmentation is a hot topic in computer vision, and hundreds of models have been proposed. For more details, please refer to the recent survey [36].

### III. METHODS

Our model consists primarily of two parts: Laplacian salience-gated filtering and pyramid feature fusion; cf. Fig. 2.

#### A. Laplacian Salience-Gated Filtering

1) *Laplacian*: The Laplacian, or Laplacian operator, is the second derivative of the  $n$ -dimensional Euclidean space, defined as

$$\Delta f = \nabla \cdot \nabla f = \sum_{i=1}^n \frac{\partial^2 f}{x_i^2}, \quad (1)$$

where  $f$  is a function of  $\mathbf{x} = \langle x_1, x_2, \dots, x_n \rangle$ . For two-dimensional space,

$$\Delta f(x, y) = \frac{\partial^2 f}{x^2} + \frac{\partial^2 f}{y^2}. \quad (2)$$

Note, the  $x$  and  $y$  are the relative distance to the interesting point  $(x_i, y_i)$ , not the absolute coordinates.

Since the second derivative is very sensitive to noise, the Laplacian is further smoothed by a Gaussian, resulting in

$$\Delta f^s(x, y) = C \Delta f(x, y) e^{-\frac{x^2+y^2}{2\sigma^2}}, \quad (3)$$

with  $\sigma$  the standard deviation of the Gaussian envelope and  $C$  the normalization factor.

2) *Laplacian Wavelet*: The Laplacian wavelet in 2D space is the Laplacian (the scaling function) modulated by a complex exponential (the wavelet function), defined as

$$g(x, y; \lambda, \phi) = \Delta f^s(x, y) \cdot e^{i(\frac{2\pi}{\lambda}(x \cos \theta + y \sin \theta) + \phi)}, \quad (4)$$

where  $\lambda$  is the wavelength and  $\phi$  is the phase offset.

The Laplacian wavelet is isotropic in its original form, but various patterns should be captured, so it is further refined as

$$\begin{aligned} g(x, y; \lambda, \phi, \gamma, \theta, \sigma) &= \Delta f^s(x', y') e^{i(\frac{2\pi}{\lambda}x' + \phi)} \\ &= C \Delta f(x', y') e^{-\frac{x'^2+y'^2}{2\sigma^2}} e^{i(\frac{2\pi}{\lambda}x' + \phi)}, \end{aligned} \quad (5)$$

where  $x' = x \cos \theta + y \sin \theta$ ,  $y' = -x \sin \theta + y \cos \theta$ ,  $\gamma$  is the spatial aspect ratio between  $x$  and  $y$ , and  $\theta$  is the rotation angle from  $(x, y)$  to  $(x', y')$ . According to the convolution theorem, the final operator applied to  $f$  can be further written as

$$g(x, y) = C(1 + \gamma - \frac{x^2 + \gamma^2 y^2}{\sigma^2}) e^{-\frac{x^2 + \gamma^2 y^2}{2\sigma^2}} e^{i(\frac{2\pi}{\lambda}x' + \phi)}. \quad (7)$$

For simplicity, the parameters are ignored. In addition, the normalization factor  $C$  can be dropped out.

3) *Laplacian Salience*: According to the Euler's formula that  $e^{ix} = \cos x + i \sin x$ , the above wavelet can be split into the real and imaginary two parts, with

$$g^R(x, y) = A \cos(2\pi \frac{x}{\lambda} + \phi) \quad (8)$$

$$g^I(x, y) = A \sin(2\pi \frac{x}{\lambda} + \phi), \quad (9)$$

where

$$A = (1 + \gamma - \frac{x^2 + \gamma^2 y^2}{\sigma^2}) e^{-\frac{x^2 + \gamma^2 y^2}{2\sigma^2}}. \quad (10)$$

Let

$$\Lambda^R(x, y) = \sum_i^K g^R(x, y; \lambda_i, \phi_i, \gamma_i, \theta_i, \sigma_i) \quad (11)$$

be the integral of  $K$  patterns, then by convolving  $\Lambda^R(x, y)$  on an image  $f(x, y)$  the specific patterns contained in  $f(x, y)$  can be fished out. Analogously, the patterns defined by  $\Lambda^I(x, y)$  can also be obtained from  $f(x, y)$  by convolution as well.

Based on the above two integrated banks of filters, the saliency map  $s$  of an image  $f$  can be obtained by

$$s(x, y) = h((f(x, y) - \bar{f}) * \Lambda^R(x, y)) + h((f(x, y) - \bar{f}) * \Lambda^I(x, y)) \quad (12)$$

where  $\bar{f}$  is the mean intensity of  $f$ ,  $*$  represents convolution operation and  $h(\cdot)$  is a normalization function.

It is known that  $\Lambda^I(x, y)$  is particularly helpful in high-lighting sudden changes, such as edges, steps and corners, because of the sign change over the 0-point. To further enhance the contrastiveness, the wavelength  $\lambda$  is optimized to  $2\sqrt{3 + \gamma\sigma}$  by solving the first-order derivative of  $\Lambda^I(x, y)$  at 0. For  $\Lambda^R(x, y)$ , it is used to smooth the saliency map so as to withstand noise and intensity variations. Similarly, its wavelength is optimized to  $4\sqrt{1 + \gamma\sigma}$ .

The normalization and subtraction are critical to the generation of saliency maps. Normalization is necessary because the optimized basic wavelets are different between the odd and even waves, while subtraction is used to remove the background so that the abnormal regions can stand out.

## B. Pyramid Feature Fusion

**1) Model Architecture:** A novel saliency-gated feature pyramid architecture is proposed to distinguish vessels from other tissues; cf. Fig 2. The feature pyramid network (FPN) [47] and its variants have been intensively explored for object detection and segmentation [48] due to its superiority in multi-scale feature fusion. However, in its original form, it is unsuitable for vessel segmentation due to the erosion, or even elimination, of small vessels during downscaling. To this end, we refine the architecture in two-fold: i) bottom-up zoom-in and top-down zoom-out to enlarge tiny vessels and obtain more detailed patterns, and ii) automatic highlighting of vessels boundaries by a high-pass filter on the saliency maps at different scales.

Let  $X \in \mathbb{R}^{w \times h \times c}$  be an image with width  $w$ , height  $h$ , and channel  $c$ . The bottom-up zoom-in of  $X$  is carried out by passing it through a ResNet50 block [49] at each reverse pyramid level. The input of the higher level is the output of the lower level with up-sampling, which scales the output two times larger. The top-down zoom-out is conducted by adding the down-sampled output of the upper level and the feature maps of the same level in the bottom-up path. See the detailed illustration in Fig 2. The zooming operation (both in and out) is repeated three times, but not more, because of the following two main reasons: i) the size of feature maps grows exponentially with zoom-in, thus easily leading to memory exhaustion if scaled too deep; ii) three times of up-scaling ( $2^3 \times 2^3 = 64$  times larger in the area) is sufficient to enlarge a tiny vessel enough for exquisite segmentation.

**2) Saliency-Gated Feature Generation:** Unlike traditional feature generation, we modulate the feature maps with gates. Let  $X^{(l)}$  be an image, either the original input or its enlarged

form, at layer  $l$ , and  $F^{(l)}$  be the feature maps produced at layer  $l$ . The gate  $G(X^{(l)})$  to be applied to  $F^{(l)}$  is calculated as

$$G(X^{(l)}) = \text{IFFT}(\text{FFT}^{\text{high-pass}}(S(X^{(l)}); \rho)), \quad (13)$$

where  $S(X^{(l)})$  is the Laplacian saliency of  $X^{(l)}$ ,  $\text{FFT}^{\text{high-pass}}(x)$  is the high-pass of fast Fourier transformation of  $x$ ,  $\text{IFFT}(x)$  is the inverse fast Fourier transformation of  $x$ , and  $\rho$  is the threshold to determine the high-pass band. Based on  $G(X^{(l)})$ , the modulated feature map of  $F^{(l)}$  is calculated by

$$F^{(l)'} = G(X^{(l)}) \odot \sigma(F^{(l)}), \quad (14)$$

where  $\odot$  is the element-wise dot product, and  $\sigma(\cdot)$  is an activation function.

Intuitively, a saliency map is helpful to focus on vessel regions as well as those ambiguous regions that resemble vessels. However, texture information is very limited and varies significantly due to the large number of tiny vessels presented. To this end, we further shape the importance of feature maps on the boundary regions, hence the high pass filter. There have two main reasons: i) the boundary regions are more consistent across different vessels regardless of their size, and ii) the boundary labels are imperfect, so more attention to them improves segmentation.

**3) Saliency-Modulated Loss:** The loss function of the model is guided by the saliency of the input image  $X$ , which is computed by

$$L = -\frac{1}{MN} \sum_i^M \sum_j^N [\alpha S(X_{ij}) Y_{ij} \ln P_{ij} + (1 - \alpha) S(X_{ij}) (1 - Y_{ij}) \ln(1 - P_{ij})], \quad (15)$$

where  $Y$  is the label,  $P$  is the prediction,  $S(X)$  is the saliency of  $X$ , and  $\alpha$  is the balancing factor within the range of 0 to 1.

Our saliency-modulated loss is ostensibly similar to Focal loss [34], but de facto different. Ours is determined from the raw input, reflecting the difficulty between vessels are other regions; while Focal loss is determined from the model, unveiling the models' distinguishability, which unfortunately can vary significantly with different inputs.

Note that, the loss modulation is slightly different from the feature gating, in which the high-pass filtering is excluded in the former one. This is because the final target is the entire vessel area, while the features are primarily based on the boundary regions.

## C. Implementation Details

The proposed model is implemented in PyTorch and validated on a cluster having four NVIDIA V100 GPUs. The learning rate is optimized by Adam [50] with an initial value of  $10^{-2}$  and a reduction factor of 0.1, while the batch size is 8.

The saliency of some liver tissues, which are very different from vessels, can be 0, so they can hardly contribute to model training. To further boost the learning of non-vessel regions, the lower bound of the saliency is set to 0.05.



## IV. EXPERIMENTS

### A. Datasets

Two types of datasets are used to evaluate the performance of our model as well as others, including the widely used public datasets and our newly constructed dataset. The public ones are 3D-IRCADb (3D Image Reconstruction for Comparison of Algorithm Database) [51] and MSD (Medical Segmentation Decathlon) [52], while ours is LiVS (short for Liver Vessel Segmentation).

The 3D-IRCADb contains 20 volumes of CT images having 2,823 slices with vessel masks, and MSD consists of 303 volumes and 21,120 slices having vessel masks.

LiVS has 532 volumes and 15,984 slices, where each slice is delineated by three senior medical imaging experts and the final mask is determined by majority vote. Due to the small size of vessels, the delineation of each vessel can oscillate easily. To address this, the coincidence of each vessel among the three masks is calculated. If the majority voting over any mask is smaller than 0.5, the vessel is highlighted and sent back to the three experts for further refinement. This process is repeated until no disagreement exists. A detailed description of each volume is available at <https://github.com/lzhLab/LiVS>.

Although SLIVER07 [53] (having 30 volumes and 4,159 slices) and CHAOS [54] (having 50 volumes and 2,874 slices) have been frequently used in liver vessel segmentation, they actually have no *vessel* mask, and all studies using these datasets have to manually delineate vessel masks. Hence, they are excluded from this study.

### B. Evaluation Metrics

The performance is measured by Dice similarity coefficient (DSC), volumetric overlap error (VOE), average symmetric surface distance (ASSD) and Hausdorff distance (HD), which are defined as:

$$DSC = 2|P \cap G| / (|P| + |G|) \quad (16)$$

$$VOE = 1 - |P \cap G| / (|P \cup G|) \quad (17)$$

$$ASSD = \frac{\sum_{x \in \partial P} d(x, \partial G) + \sum_{x \in \partial G} d(x, \partial P)}{\partial P + \partial G} \quad (18)$$

$$HD = \max\{\max_{x \in \partial P} d(x, \partial G), \max_{x \in \partial G} d(x, \partial P)\}, \quad (19)$$

where  $P$  is the prediction,  $G$  is the ground truth,  $\partial X$  is the boundary of  $X$ ,  $d(x, \partial Y)$  is the minimum Euclidean distance between  $x$  and  $\partial Y$  defined as  $d(x, \partial Y) = \min_{y \in \partial Y} d(x, y)$ .

Among them, DSC and VOE are overlap-based measurements, while ASSD and HD are boundary distance-based metrics. In practice, DSC is the most widely used one as it is more robust.

### C. Performance Qualification

The performance of our newly proposed model is examined on the three datasets, and head-to-head comparisons with existing state-of-the-art models are also carried out on the three datasets with the same sets of training, validation, and testing data as well. Two types of existing models are adopted, i.e., 2D-based and 3D-based. The typical 2D models borrowed

TABLE I

LIVER VESSEL SEGMENTATION PERFORMANCE ON 3D-IRCADb

Model	Structure	DSC $\uparrow$	VOE $\downarrow$	ASSD $\downarrow$	HD $\downarrow$
U-Net	2D	0.673	0.491	1.671	10.912
MFSNet	2D	0.572	0.598	1.925	11.832
DCSAU-Net	2D	0.672	0.490	1.682	10.620
FPN	2D	0.637	0.530	2.035	12.900
CaraNet	2D	0.571	0.598	2.724	15.370
TransFusionNet	2D	0.639	0.529	1.816	11.255
DualNet	2D	0.604	0.567	2.119	11.588
M2MRF	2D	0.607	0.563	2.004	11.415
3D U-Net	3D	0.311	0.806	4.400	23.262
V-Net	3D	0.374	0.770	2.177	11.220
Ours	2D	0.684	0.476	1.588	10.118

in this study include U-Net [13], FPN [47], MFSNet [55], DCSAU-Net [56], CaraNet [41], DualNet [45] (with backbone of HED [57] and dual-sampling modulated Dice loss), TransFusionNet [46], and M2MRF [40] (the cascade version), while the 3D models are 3D U-Net [14] and V-Net [11]. The detailed results are as follows.

1) *Performance on 3D-IRCADb*: The detailed performance is shown in Table I. It can be observed that the average performance of our model on 3D-IRCADb is 0.68, 0.48, 1.59 and 10.12 for DSC, VOE, ASSD and HD, respectively. This performance is significantly better than those generated by existing models, improving DSC by 1.63% compared to the second-best. This improvement is indeed remarkable considering the difficulty of vessel segmentation, although the absolute value looks trifling. Notably, the performance of the 2D models (U-Net, FPN, MFSNet, DCSAU-Net, CaraNet, DualNet, TransFusionNet, M2MRF and ours) is markedly better than that of the 3D models (3D U-Net and V-Net), *viz.*, greater than 0.6 versus smaller than 0.4. We speculate that this is mainly attributed to the small number of training data in 3D-IRCADb, i.e., only 20 volumes. This observation inspired us to construct a large-volume dataset.

2) *Performance on MSD*: MSD is 15 times larger than 3D-IRCADb in terms of volume; Therefore, we expect significant improvement of all models on MSD compared to those evaluated on 3D-IRCADb. Unfortunately, except for V-Net, the DSCs do not have apparent improvement, and some show even worse results; see Table II. For example, the DSC obtained from U-Net is slightly smaller than those generated from 3D-IRCADb. Regarding most of the mentioned networks and our model, they produce better results on MSDs. Noteworthy, V-Net lifts the DSC from 0.374 to 0.612, indicating that it benefits most from the large number of training data. Although our model generates the most accurate results than others, the unstable results obtained by all these models suggest that the data quality needs to be improved. To this end, we carried out comprehensive experiments on our newly constructed large-volume and high-quality data; see next.

3) *Performance on LiVS*: Promisingly, the performance of all models on LiVS has experienced a significant jump, increasing the average DSC by  $0.157 \pm 0.105$  and  $0.113 \pm 0.042$  compared to 3D-IRCADb and MSD, respectively; see details in Table III. This is obviously a tremendous advancement in liver vessel segmentation.

TABLE II  
LIVER VESSEL SEGMENTATION PERFORMANCE ON MSD

Model	Structure	DSC ↑	VOE ↓	ASSD ↓	HD ↓
U-Net	2D	0.671	0.485	2.141	12.385
FPN	2D	0.648	0.511	2.023	11.348
MFSNet	2D	0.584	0.583	2.054	11.094
DCSAU-Net	2D	0.682	0.477	1.713	10.056
CaraNet	2D	0.571	0.599	1.928	9.679
TransFusionNet	2D	0.686	0.471	1.718	10.202
DualNet	2D	0.666	0.494	1.594	9.326
M2MRF	2D	0.634	0.526	2.331	13.313
3D U-Net	3D	0.369	0.762	4.192	18.774
V-Net	3D	0.612	0.552	1.230	6.600
Ours	2D	0.699	0.457	1.427	9.097

TABLE III  
LIVER VESSEL SEGMENTATION PERFORMANCE ON LIVS

Model	Structure	DSC ↑	VOE ↓	ASSD ↓	HD ↓
U-Net	2D	0.751	0.397	0.782	3.735
FPN	2D	0.763	0.382	0.689	3.007
MFSNet	2D	0.660	0.506	0.952	4.530
DCSAU-Net	2D	0.782	0.354	0.689	3.096
CaraNet	2D	0.654	0.512	1.105	5.923
TransFusionNet	2D	0.783	0.353	0.798	3.803
DualNet	2D	0.758	0.387	0.869	4.467
M2MRF	2D	0.755	0.392	0.740	3.393
3D U-Net	3D	0.582	0.566	5.123	14.403
V-Net	3D	0.779	0.359	0.408	1.774
Ours	2D	0.803	0.329	0.552	2.363

Interestingly, V-Net performs much better than other existing models, suggesting that it is suitable for vessel segmentation from liver images provided with large-volume and high-quality input data. In addition, 3D U-Net also generates significantly better results than that of MSD, indicating that the quality of LiVS is indeed better than MSD although they have a similar scale of data volume.

As for our model, the DSC score is as high as 0.803, which is lifted by 2.6% to 38.0% in terms of DSC compared to existing models. This observation suggests that the newly designed model effectively captures actual vascular regions and suppresses other liver tissues in feature learning, even without 3D information.

#### D. Saliency Quantification

Performance qualification has demonstrated the effectiveness of the proposed Laplacian saliency. Here we quantitatively analyze the impact of the saliency on images. To this end, we calculate the ratio  $r$  of the weighted number of pixels between vessels and other liver tissues, precisely  $r = \sum_{M_{x,y}=1} w_{x,y} M_{x,y} / \sum_{M_{x,y} \neq 0} w_{x,y} (1 - M_{x,y})$ , where  $M$  is the mask of an image with  $M_{x,y} = 1$  for vessels at position  $(x, y)$ ,  $M_{x,y} = 0$  for other liver tissues, and  $M_{x,y} = -1$  for those outside the liver regions. The weight  $w_{x,y}$  at position  $(x, y)$  is 1 for every pixel before Laplacian saliency filtering, and is automatically determined by the proposed Laplacian saliency after filtering.

Fig. 3 shows the distribution of  $r$  before and after Laplacian saliency filtering. Before filtering, the averaged ratio between vessel and liver is  $0.055 \pm 0.046$ , while this value is markedly tripled to  $0.163 \pm 0.151$  after filtering. It means that the proposed Laplacian saliency filtering is effective in highlighting

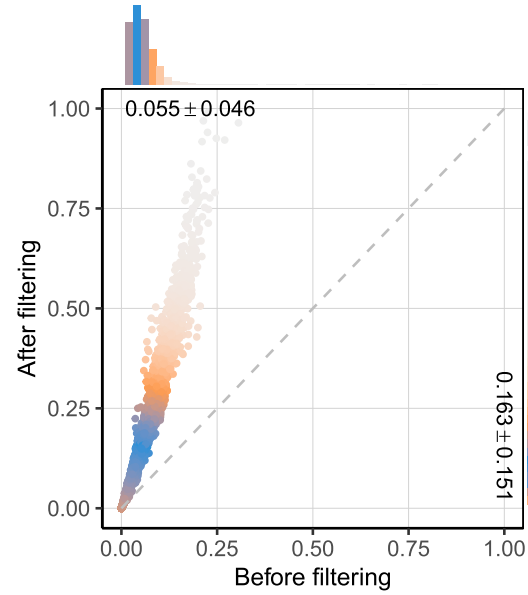


Fig. 3. Ratio between vessels and liver tissues before and after the proposed Laplacian saliency filtering.

TABLE IV  
IMPACT OF LAPLACIAN SALIENCE MODULES. EXPERIMENTS ARE CARRIED OUT ON LIVS

Model	Laplacian Saliency	High-pass Filter	DSC ↑	VOE ↓	ASSD ↓	HD ↓
FPN	✗	✗	0.763	0.382	0.689	3.007
Ours	✗	✗	0.776	0.365	0.652	3.084
Ours	✗	✓	0.765	0.378	0.758	3.556
Ours	✓	✗	0.780	0.359	0.615	2.854
Ours	✓	✓	0.803	0.329	0.552	2.363

vessel-like regions and suppressing others, which can also be observed visually in Fig. 1. Another noteworthy point is that the newly conceived filter can discriminate different classes, samples and pixels simultaneously. That being said, the imbalanced distribution of classes, the difficulty of samples, and the ambiguity of pixels can all be addressed simultaneously based on the filter.

#### E. Ablation Study

1) *Saliency Filter*: The proposed Laplacian saliency generator has two important modules, including a Laplacian saliency generator and a high-pass filter. The former is to generate the image-wise salient map highlighting vessel-like regions, while the latter is to further shape the learning on the ambiguous regions. To investigate their influence, we toggle the two modules while leaving other parts unchanged. Results show that the combination of the two yields the best performance with DSC lifted by 4% compared to the baseline. Even with the Laplacian saliency itself, the performance is still better than the original FPN model. More details can be found in Table IV.

2) *Saliency Loss*: Obviously, saliency can be incorporated into both the feature maps and the loss functions. If it is used in feature maps, the learned features will be shaped to vessels with higher attention; if it is used to modulate

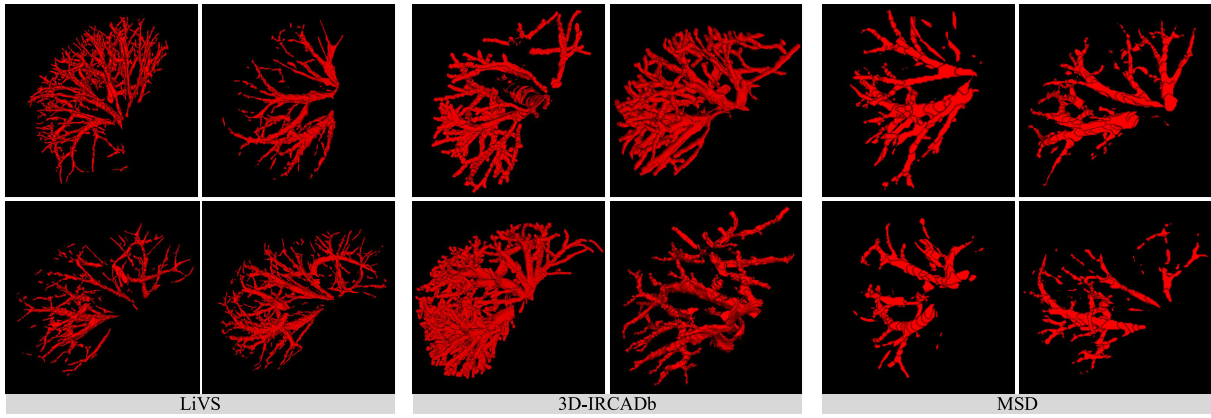


Fig. 4. Visualized 3D vessel structures built from different datasets. The density and thickness of branches produced from LiVS are evidently better than others. For 3D-IRCADb, the vessels are obviously over labeled; while MSD is under labeled on the contrary.

TABLE V

ABLATION STUDY FOR LOSS FUNCTION ON THE LIVS DATASET

Model	LS	Loss	DSC $\uparrow$	VOE $\downarrow$	ASSD $\downarrow$	HD $\downarrow$
FPN	$\times$	BCE	0.763	0.382	0.689	3.007
FPN	$\times$	SM-BCE	0.779	0.359	0.702	3.010
FPN	$\times$	Focal	0.745	0.405	0.782	3.708
Ours	$\times$	BCE	0.776	0.365	0.652	3.084
Ours	$\times$	SM-BCE	0.790	0.346	0.625	2.824
Ours	$\times$	Focal	0.767	0.376	0.740	3.707
Ours	$\checkmark$	SM-BCE	0.782	0.356	0.677	3.297
Ours	$\checkmark$	Focal	0.773	0.369	0.657	3.067
Ours	$\checkmark$	BCE	0.803	0.329	0.552	2.363

LS: Laplacian salience; SM-BCE: salience-modulated BCE loss.

the loss functions, regions with higher salience values will be more sensitive to variations. To examine which is more influential and effective, we switch on/off the modules and conduct experiments on LiVS.

Results in Table V show that the salience-gated feature together with the BCE loss achieves the best performance, while the salience-modulated loss training without the salience-gated feature gains the second best. If the model includes both the salience-gated feature and the salience-modulated loss, it unexpectedly generates slightly worse results. We speculate that the combination of salience gate and salience loss hinders the efficient learning of non-vessel features due to the very little attention paid to these regions.

Another interesting point is that the original FPN can also benefit from the use of our salience-modulated loss. This suggests that our salience is helpful in capturing vessels even if they are very small.

3) *Pyramid Level*: The feature pyramid network has proved to be effective in capturing various abstract features. However, the size of vessels is usually very small. Hence, very deep down-sampling will reduce or even eliminate vessels from the input images. To examine which level generates the best performance, we have varied the down-sampling times and conducted experiments on LiVS.

Results show that the sidelength reduced to 1/8 of the original produces the best performance. That is, the number of down-sampling pyramid levels is 3. Note, however, that the original input is first enlarged by a super-resolution layer, thus the level is increased to 5, cf. Fig. 2. Not surprisingly,

TABLE VI

PERFORMANCE OF OUR MODEL ON LIVS WITH DIFFERENT PYRAMID LEVELS

Model	Minimum sidelength	DSC $\uparrow$	VOE $\downarrow$	ASSD $\downarrow$	HD $\downarrow$
FPN	1/32	0.763	0.382	0.689	3.007
Ours	1/4	0.725	0.427	1.241	7.353
Ours	1/8	0.803	0.329	0.552	2.363
Ours	1/16	0.783	0.355	0.652	2.979
Ours	1/32	0.760	0.386	0.718	3.519

either increasing or decreasing the number of down-samplings produces worse results. For more details, please refer to Table VI.

As a comparison, the performance of the original FPN is shown in Table VI as well. It can be seen that our pyramid architecture, coupled with salience, generates much better results than FPN, even with a shallower network.

### F. Mask Examination

It is obvious that obtaining elaborated masks of vessels from liver images is very challenging because of their tiny size, heavy noise and low contrast. From the performance analysis (Section IV-C), we can see that both the volume and the quality of the datasets are critical to achieve satisfactory vessel segmentation. Precisely, LiVS has the largest number of volumes as well as the best quality, resulting in a big jump in DSC scores compared to MSD and 3D-IRCADb. Regarding the other two, they all have their own merits, with MSD having a larger volume and 3D-IRCADb having a better quality. Hence the performance of the two is comparable.

For ease of understanding, we constructed several vessel structures generated from the masks of the three datasets; see Fig. 4. Clearly, the 3D vessel structures constructed from LiVS show that the thickness of the branches is proportionate and the density of the vessels looks rational. Particularly, even skinny and tiny vessels can be visualized. Regarding 3D-IRCADb, many vessel branches are intertwined and mixed together, which is evidently impractical. For MSD, the vessel masks are heavily under-labeled, resulting in very sparse vessel structures, which is also unrealistic.

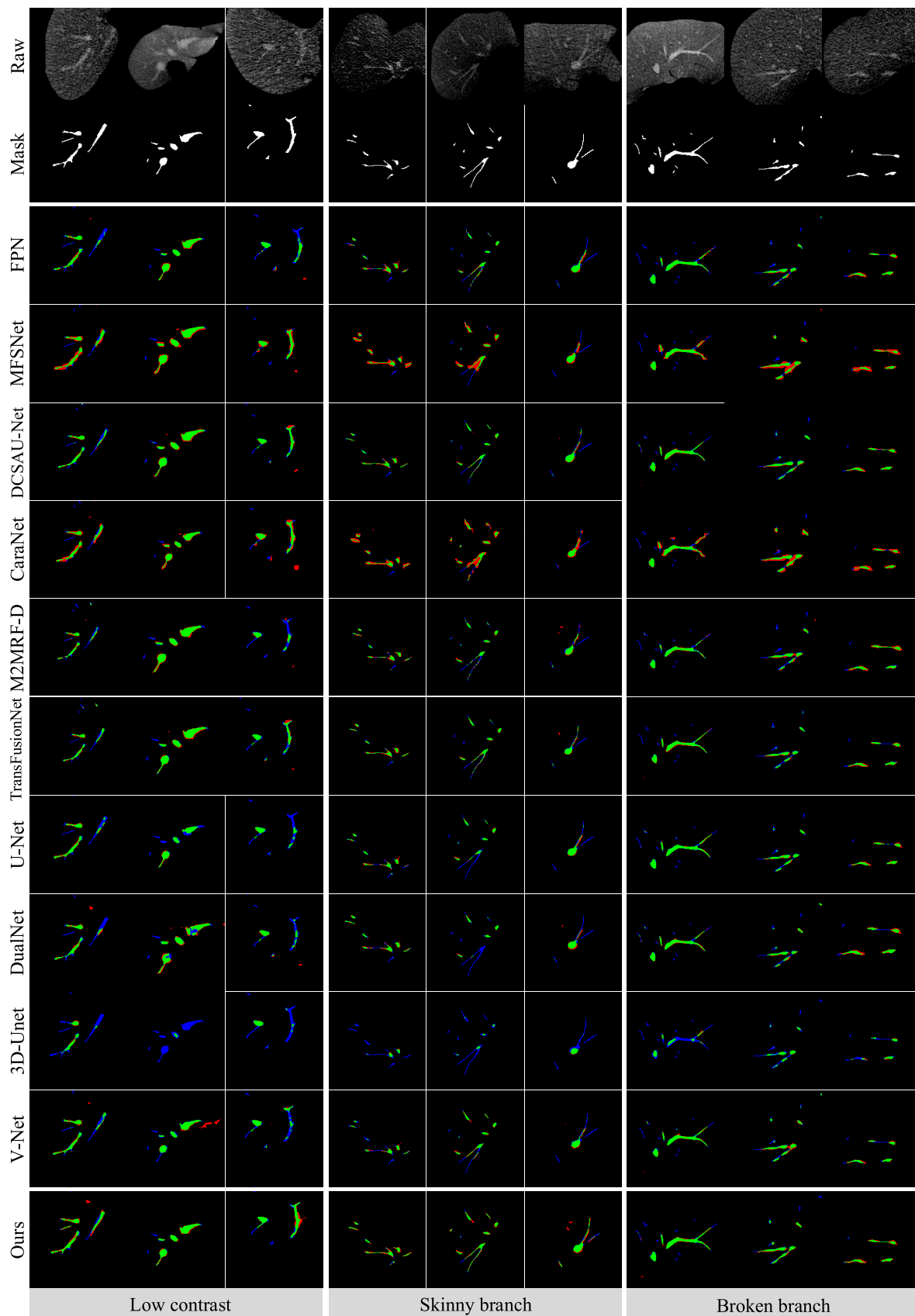


Fig. 5. Prediction visualization on LiVS. Three typical hard scenarios of vessel segmentation are presented, including vessels with low contrast, skinny branches and broken branches. The predictions are in red, while the original labels are in blue, and their superimposition is in green.



Note, this investigation is not to point out the imperfect quality of the existing or our own data. Instead, we attempt to draw attention to vessel mask purification.

### G. Case Visualization

Three main hard situations in vessel segmentation are low contrast, skinny branches and broken branches. To illustrate the performance, prediction results generated from LiVS by existing state-of-the-art models as well as ours are exemplified in Fig. 5. For vessels with low contrast, CaraNet, TransFusionNet and ours produce markedly better results than others. For those with skinny branches, only TransFusionNet and our model can generate satisfactory results, and other models miss the very skinny branches. For the broken branches, V-Net and ours perform better than the others. In short, only our newly proposed model can handle the three thorny scenarios simultaneously. This further demonstrates that the proposed Laplacian salience is sensitive to vessels and effective in vessel segmentation.

### V. CONCLUDING REMARKS

Accurate liver vessel segmentation faces two difficulties, namely the scarcity of high-quality and large-volume data as well as the sophisticated model adhering to small vessel-like regions, hence the poor performance of existing models. To this end, we have made great efforts to prepare a qualified large liver vessel dataset having 532 volumes and 15,984 slices. To investigate its goodness, a Laplacian salience-gated feature pyramid network is proposed, which can automatically focus on vessel-like regions and compensate for the skewed distribution of vessels in liver tissues. In addition, it can also put more effort into poorly discriminable regions, so that low-contrast vessels, skinny branches and broken branches can be effectively fished out. Experimental results show that the newly proposed model significantly outperforms other state-of-the-art models on both our newly proposed data and existing data. More promisingly, the Dice scores generated from our data are improved by at least 18.3% compared to those produced from other datasets. These observations suggest that our newly conceived model, as well as the data, can be of great help in advancing liver vessel segmentation.

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