



# Fully Automated Lung Lobe Segmentation in Volumetric Chest CT with 3D U-Net: Validation with Intra- and Extra-Datasets

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## Abstract

Lung lobe segmentation in chest CT has been used for the analysis of lung functions and surgical planning. However, accurate lobe segmentation is difficult as 80% of patients have incomplete and/or fake fissures. Furthermore, lung diseases such as chronic obstructive pulmonary disease (COPD) can increase the difficulty of differentiating the lobar fissures. Lobar fissures have similar intensities to those of the vessels and airway wall, which could lead to segmentation error in automated segmentation. In this study, a fully automated lung lobe segmentation method with 3D U-Net was developed and validated with internal and external datasets. The volumetric chest CT scans of 196 normal and mild-to-moderate COPD patients from three centers were obtained. Each scan was segmented using a conventional image processing method and manually corrected by an expert thoracic radiologist to create gold standards. The lobe regions in the CT images were then segmented using a 3D U-Net architecture with a deep convolutional neural network (CNN) using separate training, validation, and test datasets. In addition, 40 independent external CT images were used to evaluate the model. The segmentation results for both the conventional and deep learning methods were compared quantitatively to the gold standards using four accuracy metrics including the Dice similarity coefficient (DSC), Jaccard similarity coefficient (JSC), mean surface distance (MSD), and Hausdorff surface distance (HSD). In internal validation, the segmentation method achieved high accuracy for the DSC, JSC, MSD, and HSD ( $0.97 \pm 0.02$ ,  $0.94 \pm 0.03$ ,  $0.69 \pm 0.36$ , and  $17.12 \pm 11.07$ , respectively). In external validation, high accuracy was also obtained for the DSC, JSC, MSD, and HSD ( $0.96 \pm 0.02$ ,  $0.92 \pm 0.04$ ,  $1.31 \pm 0.56$ , and  $27.89 \pm 7.50$ , respectively). This method took  $6.49 \pm 1.19$  s and  $8.61 \pm 1.08$  s for lobe segmentation of the left and right lungs, respectively. Although various automatic lung lobe segmentation methods have been developed, it is difficult to develop a robust segmentation method. However, the deep learning-based 3D U-Net method showed reasonable segmentation accuracy and computational time. In addition, this method could be adapted and applied to severe lung diseases in a clinical workflow.

**Keywords** 3D U-net · Convolutional neural network · Deep learning · Lung lobe segmentation

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## Introduction

The human lungs are anatomically and functionally divided into five distinct compartments called lobes, which are separated by the lobar fissures. As the airway and vessel tree system are independently distributed in each lobe, many lung diseases preferentially affect a particular lobar level [1]. Emphysema, post-primary tuberculosis, and silicosis typically affect the upper lobes, whereas idiopathic pulmonary fibrosis is commonly associated with the lower lobes [2–5]. In addition, treatment options including lung cancer surgery and lung volume reduction surgery are performed at a lobar level. Therefore, lung lobe segmentation in volumetric chest computed tomography (CT) plays an important role in measuring

**Table 1** Configurations of multi-center chest CT scans at inspiration and expiration

| Chest CT dataset           | Training | Validation | Test<br>(internal dataset) | Test<br>(external dataset) | Total |
|----------------------------|----------|------------|----------------------------|----------------------------|-------|
| Asan Medical Center        | 172      | 24         | 20                         | –                          | 216   |
| Kangbuk Samsung Hospital   | 100      | 16         | 20                         | –                          | 136   |
| Konkuk University Hospital | –        | –          | –                          | 40                         | 40    |
| Total                      | 272      | 40         | 40                         | 40                         | 392   |

lung functions and regional image analysis including texture and shape analyses at a lobar level and surgical planning. Manual lobe segmentation is too time-consuming and involves some degree of inter-observer variability; thus, fully automated lung lobe segmentation is needed. The lobar fissures, which are the basis for lobe segmentation, have similar intensities to those of their neighboring structures such as pulmonary vessels and are anatomically different between individuals; fissures are often incomplete or absent and can vary greatly in shape and location, especially in the presence of a disease (e.g., infiltrative lung disease). These characteristics could lead to segmentation error in automated segmentation methods.

Various automatic approaches have been proposed for lung lobe segmentation. Lassen et al. proposed a method based on Hessian analysis to model sheet-like, tubular, and blob-like structures by the eigenvalues of the Hessian matrix, which is a matrix built of the second partial derivatives of an image [6, 7]. This method worked well in normal cases but was sensitive to CT scan protocols/parameters, image noise, and incompleteness of fissures; in particular, it did not work in the case of chronic obstructive pulmonary disease (COPD). Zhang et al. presented an atlas-based method that trains the structural shapes based on existing gold standards [8]. This method achieved good results but was time-consuming and it showed a relatively poorer performance when the disease changed the structures of the lungs. There are other segmentation methods based on thresholding, surface fitting [9], watershed [10], and graph searching with shape constraints [11]. However, these methods need parameter optimization (e.g., smoothness

and thresholding values) for the best performance because they are vulnerable to the image quality level. In addition, intensity-based methods are challenging due to the similar intensity to those of image noise, vessels, airway walls, and other structures. Therefore, these fully automated segmentation methods could not be used in clinical applications.

Recently, deep learning architectures have been successfully established for lung lobe segmentation. George et al. used a progressive, holistically nested network [12] to segment lobar boundaries, generating seed points and edge probabilities for a random walker, which could produce a final lobe segmentation mask [13]. This method showed good results not only for the normal group but also for the COPD group. However, it relied on prior segmentation of the lobar boundaries (i.e., fissures) and thus did not work well for lungs with fake or incomplete fissures.

In this study, we performed lung lobe segmentation using a deep learning technique to develop a robust algorithm without lobar fissure detection. We applied a 3D convolutional neural network (CNN) to exploit the geometric information of the lungs. Our method outperformed image processing-based segmentation in terms of accuracy and execution time. Validation with internal and external datasets demonstrated that our fully automated segmentation method could be applied to clinical radiology.

## Materials and Methods

### Patient Datasets

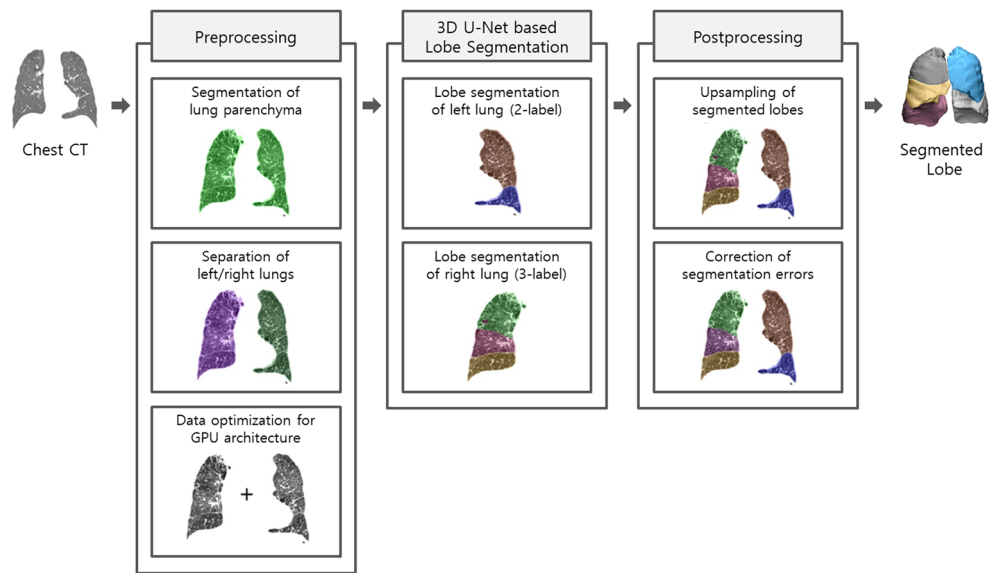
For algorithm development, we obtained the chest CT scans of mild-to-moderate COPD patients at full inspiration and expiration, which included three different datasets from Kangbuk Samsung Hospital (68 patients), Asan Medical Center (108 patients), and Konkuk University Hospital (20 patients) in South Korea. These multi-center datasets were composed of independent heterogeneous sets with different scanners, reconstruction kernels, and severities of COPD: (1) Asan Medical Center—a 16-channel multi-detector CT (MDCT) scanner (Somatom Sensation 16; Siemens Healthcare, Erlangen, Germany) with 140 kVp, 100 eff. mA/s, a slice thickness of 0.75 mm, and a reconstruction kernel of B30f; (2) Kangbuk Samsung Hospital—a 40-channel MDCT scanner (Brilliance 40; Philips Healthcare, Best, The Netherlands)

**Table 2** Basic conditions for each localized structure

| $\lambda_1$ | $\lambda_2$ | $\lambda_3$ | Structure orientation          |
|-------------|-------------|-------------|--------------------------------|
| L           | L           | L           | Noise (no preferred structure) |
| L           | L           | H–          | Bright sheet-like structure    |
| L           | L           | H+          | Dark sheet-like structure      |
| L           | H–          | H–          | Bright tubular structure       |
| L           | H+          | H+          | Dark tubular structure         |
| H–          | H–          | H–          | Bright blob-like structure     |
| H+          | H+          | H+          | Dark blob-like structure       |

$|\lambda_1| \leq |\lambda_2| \leq |\lambda_3|$  are eigenvalues of Hessian matrix, L:  $\lambda_i \cong 0$ , H+:  $\lambda_i$  is high positive value, H–:  $\lambda_i$  is high negative value

**Fig. 1** Overall process of the proposed lung lobe segmentation method



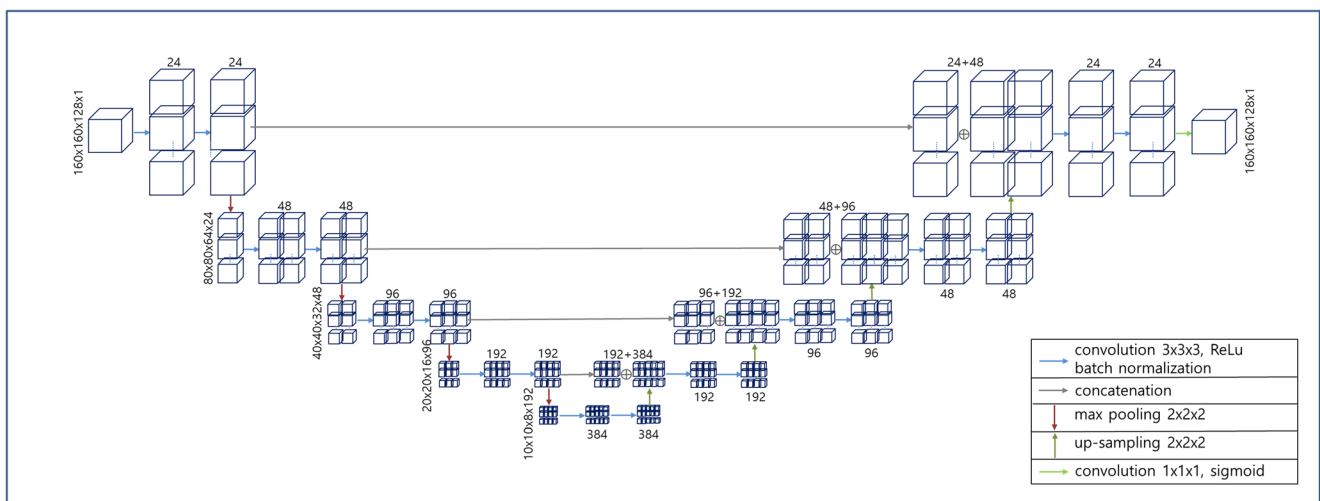
with 140 kVp, 110 mA/s, a slice thickness of 0.8 mm, and a reconstruction kernel of B; (3) Konkuk University Hospital—a 64-channel MDCT scanner (Brilliance 64; Philips Healthcare, Best, The Netherlands) with 140 kVp, 183 mA/s, a slice thickness of 0.625 mm, and a reconstruction kernel of B. These datasets were split into training (272 CT scans), validation (40 scans), and test (40 scans) datasets. The CT scans of Konkuk University Hospital were used only for external validation (see Table 1).

**Gold Standard of Lung Lobe Segmentation**

Since manual lobe segmentation is too time-consuming and difficult, we semi-automatically segmented all images based on Hessian analysis [6], which was previously developed. The Hessian analysis is a segmentation technique based on the Hessian matrix which is derived from the second derivatives:

$$H(x, y, z) = \nabla^2 \begin{bmatrix} f_{xx} & f_{xy} & f_{xz} \\ f_{yx} & f_{yy} & f_{yz} \\ f_{zx} & f_{zy} & f_{zz} \end{bmatrix} \tag{1}$$

The eigenvalues ( $|\lambda_1| \leq |\lambda_2| \leq |\lambda_3|$ ) of the Hessian matrix characterize the local morphological structure of an object, such as sheet-like, tubular, and blob-like structures [6] (see Table 2). The lobar fissures were segmented using the conditions of sheet-like structures. To characterize the structures of interest, suppress local noises and perform Hessian analysis in CT images, we used Gaussian smoothing with  $3 \times 3 \times 3$  kernel convolution before  $3 \times 3 \times 3$  Hessian analysis with 1-pixel stride moving window. These segmented lobe regions were confirmed and modified by expert radiographers with more than 5 years of experience using the in-house software AView, which included several modules of lung segmentation, left/right lung separation, Hessian analysis-based lobe segmentation, and manual editing tools.



**Fig. 2** Architecture of 3D U-Net

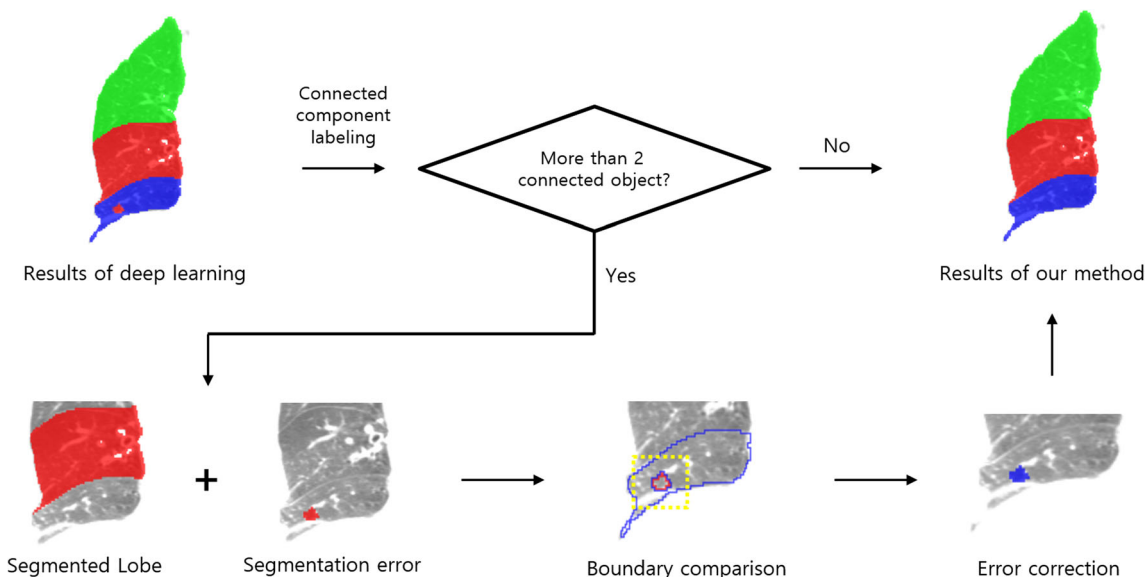


Fig. 3 Process of segmentation error correction

### Fully Automated Lung Lobe Segmentation Using 3D U-Net

Lung lobe segmentation would likely be better performed by 3D CNN, since lung lobes are 3D objects, and 2D CNN do not consider 3D continuity between slices, which guarantees 3D continuity of segmented lobes. In addition, since it is frequently found that human lungs have incomplete and/or fake fissures [14], fissure-based lung lobe segmentation could be susceptible to these anatomical variations. We, therefore, introduced 3D CNN to fully automated 3D lung lobe segmentation without lobar fissure detection. Our system proceeds through three main steps (Fig. 1). To segment the left and right lung lobes with different geometric structures, we split whole lungs into the left and right lungs and optimize them for GPU architectures. From each of left and right lungs, the lobes are segmented using 3D U-Net. Finally, the segmentation errors are corrected by boundary comparison.

### Preprocessing

The geometric structures of the left and right lungs are different (not symmetrical) as the left lung has two lobes but the right lung has three. Our method trained these different left and right lungs separately. First, we segmented the lungs by thresholding (ranging from  $-1024$  HU to  $-400$  HU) and using region-growing techniques [15]. To handle the hilar regions, the airways were excluded by thresholding (ranging from  $-1024$  HU to  $-950$  HU) and region growing with an initial seed point, which was automatically detected by the circle detection algorithm. As the airways have circle-like shapes in the axial plane, they could be excluded by the modified Hough transform [16], which was performed on the upper slices (5%) based on the z-axis. Second, the segmented lungs were split into the left and right lungs. When the left and right lungs were weakly connected, they were easily separated using a small number of iterations based on 3D morphological operations, 3D distance transform, and surface fitting algorithms [17, 18]. To check if the lungs are

Table 3 Definitions of evaluation metrics

| Evaluation metric | Definition   |
|-------------------|--|
| DSC ( $X, Y$ )    | $\frac{2 X \cap Y }{ X  +  Y }$ [2]  |
| JSC ( $X, Y$ )    | $\frac{ X \cap Y }{ X \cup Y }$ [3]  |
| MSD ( $X, Y$ )    | $\frac{1}{N_x + N_y} \left\{ \sum_{x \in S_x} \min_{y \in S_y} (\text{dist}(x, y)) + \sum_{y \in S_y} \min_{x \in S_x} (\text{dist}(x, y)) \right\}$ [4] |
| HSD ( $X, Y$ )    | $\max \left\{ \sup_{x \in S_y} \inf_{y \in S_y} \text{dist}(x, y), \sup_{y \in S_y} \inf_{x \in S_x} \text{dist}(x, y) \right\}$ [5]                     |

Optimal value of DSC, JSC, MSD, and HSD was 1, 1, 0, and 0, respectively

**Table 4** Performance of the proposed lung lobe segmentation method evaluated with the internal-dataset

| <i>N</i> = 40 | LtLower        | LtUpper        | RtLower        | RtMiddle       | RtUpper        | Overall        |
|---------------|----------------|----------------|----------------|----------------|----------------|----------------|
| DSC           | 0.9721 ± 0.011 | 0.9806 ± 0.007 | 0.9684 ± 0.012 | 0.9436 ± 0.019 | 0.9753 ± 0.012 | 0.9680 ± 0.018 |
| JSC           | 0.9460 ± 0.021 | 0.9620 ± 0.014 | 0.9389 ± 0.022 | 0.8938 ± 0.033 | 0.9520 ± 0.022 | 0.9385 ± 0.031 |
| MSD (mm)      | 0.5796 ± 0.240 | 0.5092 ± 0.231 | 0.6901 ± 0.246 | 1.0239 ± 0.415 | 0.6545 ± 0.413 | 0.6914 ± 0.364 |
| HSD (mm)      | 14.40 ± 5.85   | 18.20 ± 12.06  | 17.59 ± 13.68  | 18.55 ± 8.47   | 16.86 ± 13.24  | 17.12 ± 11.07  |

*Lt*, left; *Rt*, right

separated, connected component labeling [19] were used. At this stage, if only one connected object was detected, we judged this lung has strongly connected, the next step was performed. When the left and right lungs were strongly connected, the junction parts between them were weakened by morphological operation and Hessian analysis. In a similar manner as in the generation of the gold standard, the junction parts were detected and excluded using a combination of eigenvalues for sheet-like structures. Finally, the separated left/right lungs were resized to 160×160×128 in order to optimize the data for GPU architectures. In our method, lung segmentation and half-lung separation of our previous study were performed to obtain rough coordinates and size of square-cropped each lung image [17].

**Generalized Segmentation Using 3D U-Net**

In order to capture the 3D information of the lobar fissures, we applied 3D U-Net [20], which extended the U-Net architecture from Ronneberger et al. [21] by replacing all 2D operations with their 3D counterparts. U-Net, which is a type of fully convolutional network [22], is one of the most widely used CNN architectures for image segmentation. 3D U-Net consists of the analysis and synthesis paths, and each path has four resolution layers (Fig. 2). In the analysis path, each layer is composed of two steps of 3×3×3 convolution followed by a rectified linear unit (ReLU) as well as 2×2×2 max pooling with strides of two in each dimension, which can reduce dimensions. In the synthesis path, each layer is composed of an up-convolution of 2×2×2 by strides of two in each dimension as well as two 3×3×3 convolutions followed by a ReLU,

which can extend dimensions. One of the advantages of 3D U-Net is the concatenation of layers of equal resolution, preventing loss of information at the deeper layers.

We applied the 3D U-Net architecture to lung lobe segmentation with the structural hyper-parameters including an input size of 160×160×128, since the size of input images needs to be unified for training. Our 3D U-Net was implemented in Keras using Theano, and we used cuDNN convolution layers for efficient memory management. Given the training datasets (272 cases for training and 40 cases for validation), our 3D U-Net was learned by minimizing the training error, i.e., the Dice similarity coefficient [23] (DSC) between the inferred label and the gold standard label. The training error was minimized by running the back-propagation algorithm and it took around 3 days using TITAN XP with 12 GB GPU. With separately trained optimal models for the left and right lungs, we then segmented the lung lobes. The final lung lobes were obtained by upsampling and correction of segmentation error. Since each lung lobe is a single continuous object, we defined that they included the segmentation error in case of the segmentation results having more than two objects. These problems sometimes happened, and we detected the segmentation errors using connected component labeling, and then these errors were corrected by boundary comparison (Fig. 3).

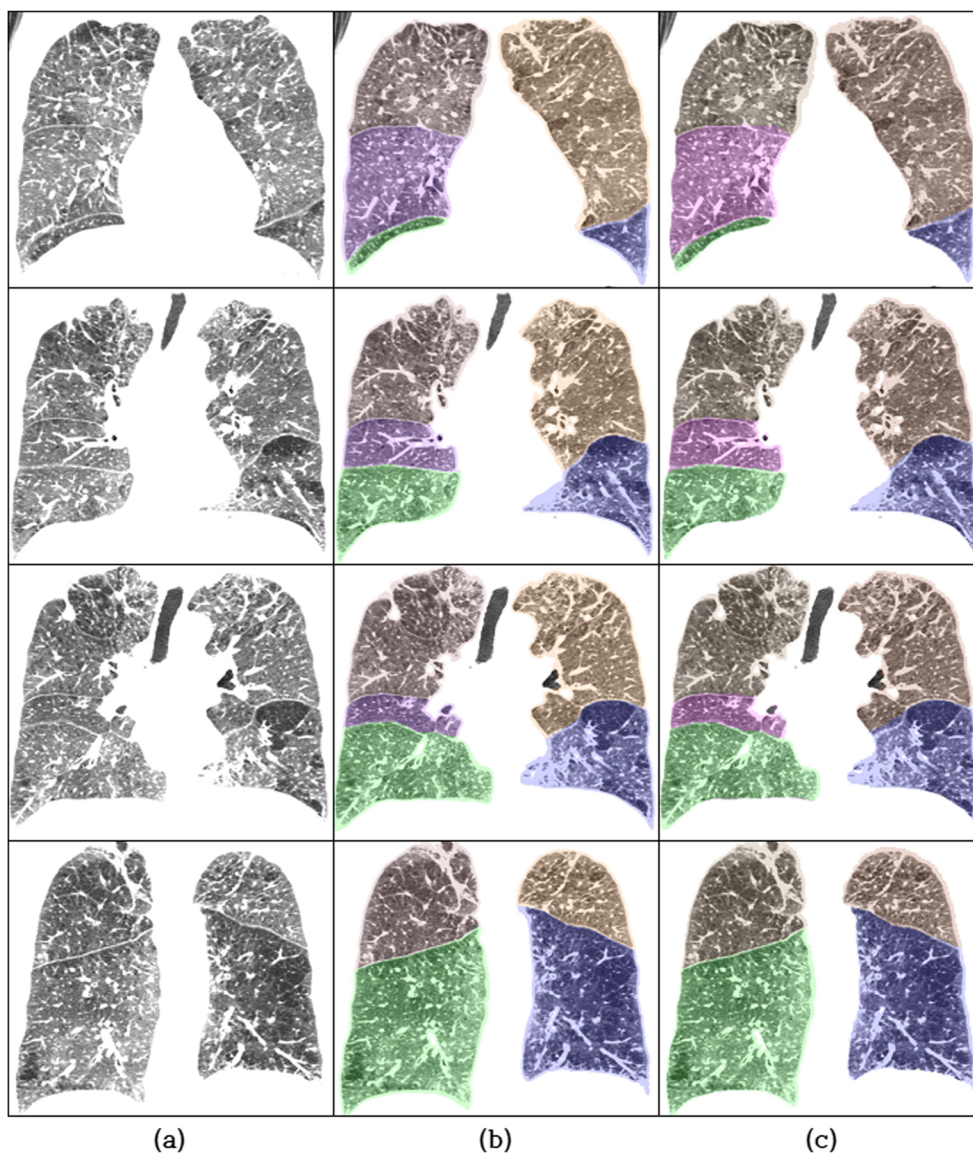
**Evaluation Metric and Statistical Analysis**

In this study, three methods including Hessian analysis-based segmentation [6] with manual correction, semi-automatic segmentation with manual correction using the in-house software

**Table 5** Comparison of the performance and execution time of different methods

| <i>N</i> = 5           | Hessian analysis-based segmentation (with manual correction) | Grid-fitting-based segmentation (with manual correction) | Our method (fully automated) |
|------------------------|--|--|------------------------------|
| Left lung (s)          | 220.12 ± 66.20   | 83.00 ± 13.87  | 6.49 ± 1.19                  |
| Right lung (s)         | 357.22 ± 90.02   | 200.60 ± 43.75   | 8.61 ± 1.08                  |
| Accuracy (overall DSC) | 0.915 ± 0.10   | 0.962 ± 0.022  | 0.948 ± 0.018                |
| Accuracy (overall JSC) | 0.857 ± 0.150  | 0.928 ± 0.040  | 0.901 ± 0.032                |
| Accuracy (overall MSD) | 2.48 ± 3.22  | 0.90 ± 0.46  | 1.36 ± 0.75                  |
| Accuracy (overall HSD) | 45.23 ± 28.80  | 21.82 ± 7.31   | 28.93 ± 14.24                |

**Fig. 4** Result images of the external validation. Rows from upper to lower show coronal images with segmentation results from anterior to posterior of a patient CT scan. **a** Chest CT scans. **b** Gold standard. **c** Segmentation results



AView, and our proposed method were compared in terms of segmentation accuracy and processing time. All of the three methods assumed that the lung segmentation and separation of the left and right lungs are complete. Hessian analysis-based segmentation requires user interaction as it is highly sensitive to parameter settings and contains false positives. In the case of semi-automatic segmentation, we used a grid-fitting algorithm [24] including a manual drawing of fissures and corrections.

The performance of the proposed method was evaluated with two types of independent datasets: internal and external datasets. Each segmentation result was evaluated and compared with the gold standard. To quantify the comparisons, we computed the DSC and Jaccard similarity coefficient [25] (JSC). In addition, the mean surface distance (MSD) and Hausdorff surface distance [26] (HSD) were also computed as the DSC and JSC are presented as a high value in the case of large objects. Our evaluation metrics are defined in

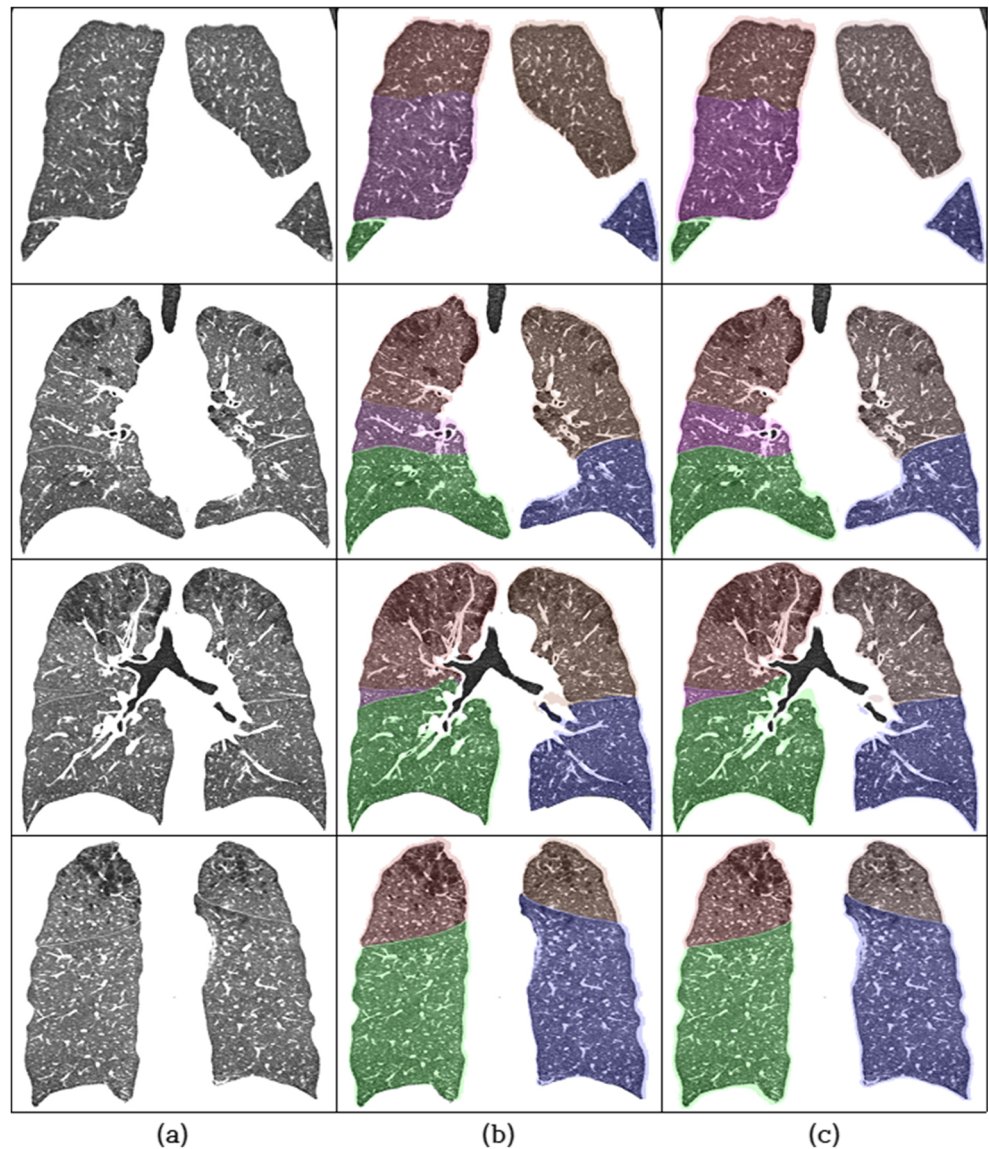
Table 3.  $|X|$  is the number of voxels in volume  $X$ , and  $S_X$  is the surface (i.e., border voxels) of  $X$ .  $N_X$  is the number of voxels on  $S_X$ , and  $\text{dist}(x, y)$  is the voxel distance between  $x$  and  $y$ .

## Results

### Internal Validation of Lung Lobe Segmentation

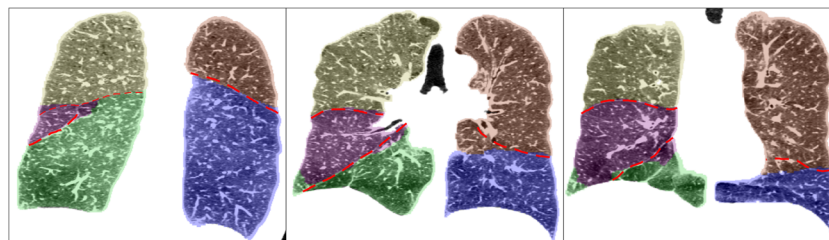
The analysis of our datasets was challenging as they were derived from mild-to-moderate COPD patients. Table 4 summarizes the performance of all four metrics (DSC, JSC, MSD, and HSD) with the internal dataset (40 cases not used for training). The overall DSC, JSC, MSD, and HSD were  $0.9680 \pm 0.018$ ,  $0.9385 \pm 0.031$ ,  $0.6914 \pm 0.364$ , and  $17.12 \pm 11.07$ , respectively. Although the segmentation of the right middle lobe is usually difficult, we obtained good

**Fig. 5** Result images of external validation in case of incomplete fissures. Rows from upper to lower show coronal images with segmentation results from anterior to posterior of a patient CT scan. **a** Chest CT scans. **b** Gold standards. **c** Segmentation results



results with a DSC of  $0.9436 \pm 0.019$  and JSC of  $0.8938 \pm 0.033$ . Table 5 shows a comparison of the average execution time of three methods including Hessian analysis-based segmentation with manual correction, semi-automatic segmentation with manual correction using the in-house software AView, and our proposed method. The Hessian analysis-based segmentation and semi-automatic segmentation

methods required manual corrections or interactions; thus, the execution time was long. Although our method could be used with a large number of datasets, we used only five cases for comparison with other methods; these five cases were randomly selected from the internal datasets. These experiments were performed for comparing execution time of each method without algorithms failure. Our fully automated



**Fig. 6** Segmentation results of coronal images of a poor case with fissures of the gold standard (red dashed lines) from anterior to posterior (left to right images). Quantification results including DSC, JSC, MSD, and HSD were 0.9445, 0.8954, 1.698, and 31.482, respectively

method required an average of  $6.49 \pm 1.19$  s for the left lobes and  $8.61 \pm 1.08$  s for the right lobes on Intel Xeon E5-2620 v3 @ 2.4 GHz, 6-core CPU, 32 GB RAM, GeForce GTX 1080 with 8 GB, and Windows 10 OS. Figures 4 and 5 show that our method was robust even in cases with incomplete fissures, while Fig. 6 shows the case of poor performance.

### External Validation of Lung Lobe Segmentation

In order to evaluate the generalizability of our model, we performed an external validation using an independent dataset with the training step. The external dataset was from Konkuk University Hospital, which was scanned with a different scanner and reconstruction kernel. Table 6 summarizes the performance of the external validation with all four metrics (DSC, JSC, MSD, and HSD). Although this dataset was substantially different from the training dataset, our method achieved a high accuracy with a DSC of  $0.96 \pm 0.02$ , JSC of  $0.92 \pm 0.03$ , MSD of  $1.31 \pm 0.56$ , and HSD of  $27.89 \pm 7.50$ . Especially for the right middle lobe, which is a very challenging task for segmentation that shows a reasonable accuracy with a DSC of  $0.93 \pm 0.03$ , JSC of  $0.87 \pm 0.05$ , MSD of  $1.48 \pm 0.84$ , and HSD of  $23.29 \pm 8.94$ . In general, our method shows a robust performance on all test cases with higher accuracies with small standard deviations. Furthermore, we evaluated the performance of our method according to COPD severity based on COPD GOLD guidelines [27]. The higher COPD GOLD stage means more severe COPD. Table 7 shows good and similar results regardless of COPD severity.

### Discussion

Previously proposed automatic techniques for lung lobe segmentation are sensitive to anatomic and disease variations, reconstruction kernels, and parameter settings, making it impossible to fully automate the segmentation of the data. These techniques could be used only for the initial segmentation with manual correction and parameter settings. 3D lung lobe segmentation is costly, time-consuming, and laborious; it frequently requires manual correction by expert radiologists, which is a barrier to application in a clinical workflow. To

overcome these problems, we proposed a fully automated lung lobe segmentation method, which could produce high-quality results without manual correction. George et al. [12] proposed another method of lung lobe segmentation using progressive holistically nested networks and the random walker algorithm. The method in this study also demonstrated good results for the normal group and the COPD group. However, similar to most other methods, it relied on prior segmentation of the lobar fissures, which had problems with fake or incomplete fissures. Fake fissures could generate segmentation error and negatively affect the segmentation result. In addition, the method would not work in the case of incomplete fissures. Some automatic algorithms produce less accurate results because they could not distinguish minor fissures on the right lung. To address these problems, we trained the lung lobe without lobar fissure information (Fig. 6).

Parameter optimization of the conventional methods for lung lobe segmentation is very sensitive to disease severity. The evaluation of our method according to the COPD GOLD stage demonstrated the robustness of our method to disease variation.

In this study, there are several limitations. We selected mild-to-moderate COPD patients in this study. However, when a lung disease becomes severe, the lung structures would change considerably. For a more robust clinical outcome, patients with strong anatomical variations, various lung diseases including severe COPD with emphysema, diffuse infiltrative lung disease (DILD) with complex parenchymal texture patterns, post-primary tuberculosis, and cancer which could have potentials to change pathological lung and lobe anatomy should be included. In addition, quality of lung segmentation could affect that of lobe segmentation, especially in DILD patients, which could be enhanced by developing deep learning method for lung segmentation. In future studies, we will evaluate the performance of this method using open dataset on lung lobe segmentation and develop a more robust lung segmentation method based on this method using more high-quality segmentation labels. In addition, we may build larger datasets for lung lobe segmentation, which could lead to the development of another robust lobe segmentation method. To build larger datasets, we could develop an easy user interface, fast processing, initial segmentation techniques, and correction modules.

**Table 6** Performance of the proposed lung lobe segmentation method evaluated with the external dataset

| N = 40   | LtLower            | LtUpper            | RtLower            | RtMiddle           | RtUpper            | Overall            |
|----------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|
| DSC      | $0.9556 \pm 0.013$ | $0.9701 \pm 0.010$ | $0.9697 \pm 0.007$ | $0.9306 \pm 0.030$ | $0.9697 \pm 0.007$ | $0.9561 \pm 0.022$ |
| JSC      | $0.9152 \pm 0.024$ | $0.9423 \pm 0.018$ | $0.9413 \pm 0.013$ | $0.8716 \pm 0.049$ | $0.9413 \pm 0.013$ | $0.9167 \pm 0.031$ |
| MSD (mm) | $1.3762 \pm 0.504$ | $1.0625 \pm 0.440$ | $1.1591 \pm 0.355$ | $1.4804 \pm 0.841$ | $1.1591 \pm 0.355$ | $1.3149 \pm 0.563$ |
| HSD (mm) | $29.03 \pm 6.61$   | $27.74 \pm 7.54$   | $30.83 \pm 5.24$   | $23.29 \pm 8.94$   | $30.83 \pm 5.24$   | $27.89 \pm 7.50$   |

Lt, left; Rt, right



**Table 7** Performance of the proposed lung lobe segmentation method according to COPD GOLD stage level

|                          |          | LtLower        | LtUpper        | RtLower        | RtMiddle       | RtUpper        | Overall        |
|--------------------------|----------|----------------|----------------|----------------|----------------|----------------|----------------|
| Normal (N=2)             | DSC      | 0.9542 ± 0.014 | 0.9711 ± 0.007 | 0.9425 ± 0.014 | 0.9411 ± 0.013 | 0.9676 ± 0.007 | 0.9553 ± 0.016 |
|                          | JSC      | 0.9125 ± 0.025 | 0.9440 ± 0.013 | 0.8913 ± 0.026 | 0.8889 ± 0.028 | 0.9372 ± 0.014 | 0.9148 ± 0.029 |
|                          | MSD (mm) | 0.9613 ± 0.120 | 0.7815 ± 0.108 | 1.1452 ± 0.079 | 1.1119 ± 0.142 | 1.1061 ± 0.116 | 1.0212 ± 0.167 |
| COPD GOLD stage 1 (N=8)  | HSD (mm) | 22.11 ± 2.82   | 21.99 ± 1.16   | 19.53 ± 1.22   | 17.40 ± 1.59   | 26.95 ± 3.45   | 21.60 ± 3.76   |
|                          | DSC      | 0.9464 ± 0.017 | 0.9623 ± 0.016 | 0.9594 ± 0.011 | 0.9359 ± 0.012 | 0.9683 ± 0.008 | 0.9544 ± 0.017 |
|                          | JSC      | 0.8986 ± 0.031 | 0.9277 ± 0.029 | 0.9221 ± 0.020 | 0.8798 ± 0.021 | 0.9386 ± 0.014 | 0.9133 ± 0.031 |
| COPD GOLD stage 2 (N=18) | MSD (mm) | 1.9134 ± 0.802 | 1.5553 ± 0.754 | 1.7244 ± 0.780 | 1.3699 ± 0.314 | 1.4257 ± 0.658 | 1.5977 ± 0.680 |
|                          | HSD (mm) | 31.36 ± 8.15   | 28.60 ± 9.19   | 29.12 ± 7.15   | 24.01 ± 7.93   | 29.54 ± 5.59   | 28.53 ± 7.70   |
|                          | DSC      | 0.9560 ± 0.011 | 0.9716 ± 0.007 | 0.9529 ± 0.014 | 0.9339 ± 0.012 | 0.9681 ± 0.007 | 0.9565 ± 0.017 |
| COPD GOLD stage 3 (N=12) | JSC      | 0.9159 ± 0.021 | 0.9448 ± 0.013 | 0.9103 ± 0.026 | 0.8762 ± 0.022 | 0.9382 ± 1.233 | 0.9171 ± 0.031 |
|                          | MSD (mm) | 1.2691 ± 0.323 | 0.9111 ± 0.210 | 1.4455 ± 0.287 | 1.3495 ± 0.457 | 1.0423 ± 0.147 | 1.2035 ± 0.357 |
|                          | HSD (mm) | 28.00 ± 7.21   | 24.94 ± 4.42   | 29.11 ± 8.01   | 20.89 ± 6.21   | 28.61 ± 4.16   | 26.31 ± 6.79   |
|                          | DSC      | 0.9612 ± 0.010 | 0.9732 ± 0.006 | 0.9556 ± 0.013 | 0.9205 ± 0.052 | 0.9735 ± 0.006 | 0.9568 ± 0.031 |
|                          | JSC      | 0.9255 ± 0.018 | 0.9479 ± 0.012 | 0.9153 ± 0.024 | 0.8565 ± 0.083 | 0.9484 ± 0.012 | 0.9187 ± 0.052 |
|                          | MSD (mm) | 1.2478 ± 0.225 | 1.0080 ± 0.149 | 1.4789 ± 0.256 | 1.8194 ± 1.392 | 1.1653 ± 0.251 | 1.3424 ± 0.691 |
|                          | HSD (mm) | 30.19 ± 3.99   | 32.34 ± 8.68   | 28.74 ± 4.16   | 27.39 ± 12.21  | 35.68 ± 3.42   | 30.87 ± 7.67   |

## Conclusion

In this study, we proposed a fully automated lung lobe segmentation method using U-Net. Since U-Net shows a good performance on image segmentation, we customized the U-Net into 3D and optimized it for lung lobe segmentation in 12-bit medical images with memory limitation of current GPU. In addition, we designed an overall process from preprocessing to error correction, which leads the lung lobe segmentation with high accuracy.

Our method was highly effective not only for the normal group but also for the mild COPD group. Furthermore, it performed well in the presence of fake and incomplete fissures as the method did not rely on the information of the lobar fissures. Our method could be used without manual correction, and we markedly improved the execution time compared with that of other image processing-based methods. To address the overfitting problem of typical deep learning methods, we trained our model using two different datasets, which included independent heterogeneous sets with different scanners and reconstruction kernels. We performed an external validation to evaluate the generalizability of our model.

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