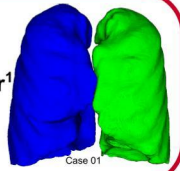




# Automatic Lung Segmentation in MDCT Images

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## Introduction

A fast and reliable segmentation of the lung tissues is of fundamental importance for many clinical applications like emphysema quantification or lung nodule detection. We present a method for fully automated extraction of the lungs from volumetric computed tomography (CT) images. The method is part of our in-house software *yacta*.

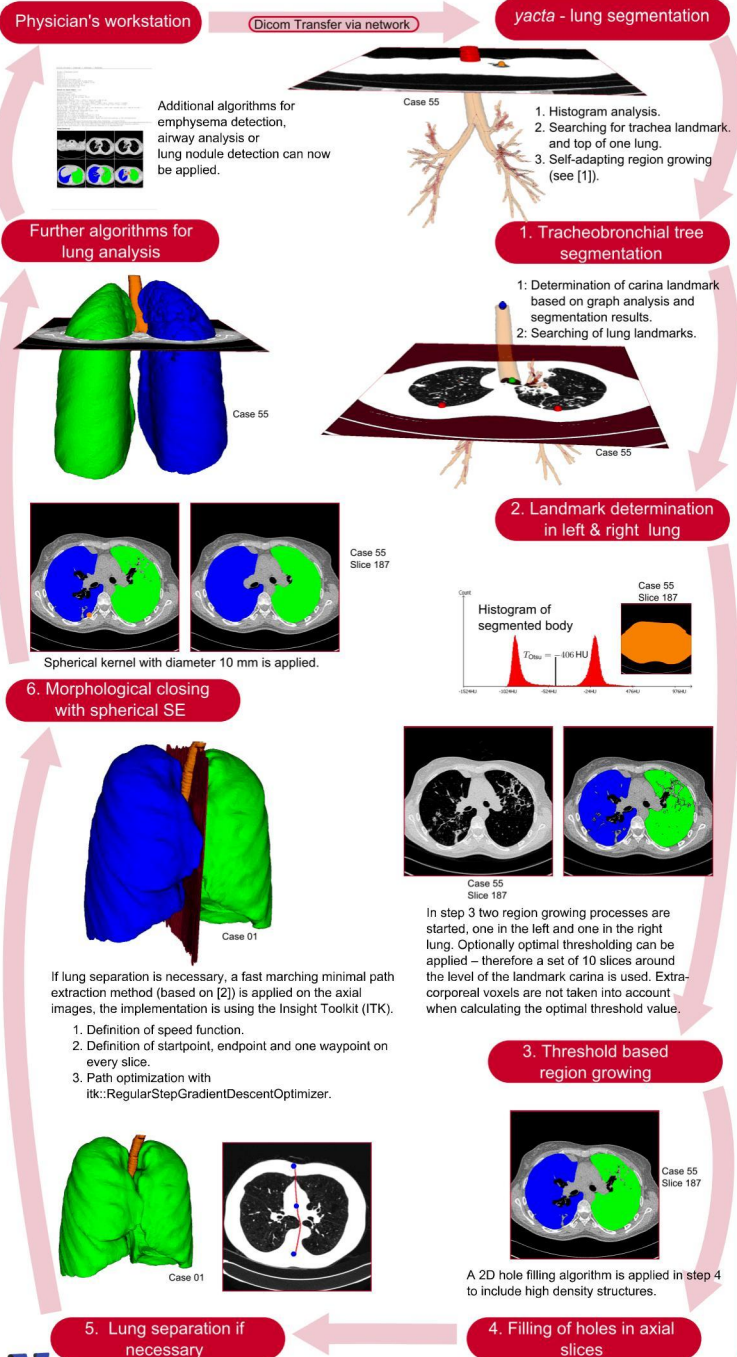
Depending on anatomical variations and pathological changes lung segmentation can be demanding. On the one hand clinical applications should be able to handle severe pathologic lung CT images, e.g. for longitudinal studies. On the other hand they should also be sensitive enough to find pathologic lung changes in an early state because a precocious diagnosis is the best way to ensure the success of a treatment.

An evaluation on 55 clinical datasets (LOLA11-datasets) shows that the segmentations of our method achieves a good average volumetric overlap (97%) against manually generated reference segmentations.

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The algorithm currently used in *yacta* can be classified as an advanced conventional method and consists of following steps:

1. Segmentation of tracheobronchial tree.
2. Landmark determination in left and right lung.
3. Threshold based region growing.
4. Filling of holes in axial slices.
5. Lung separation if necessary.
6. Morphological closing operation with a spherical structural element.



## Results

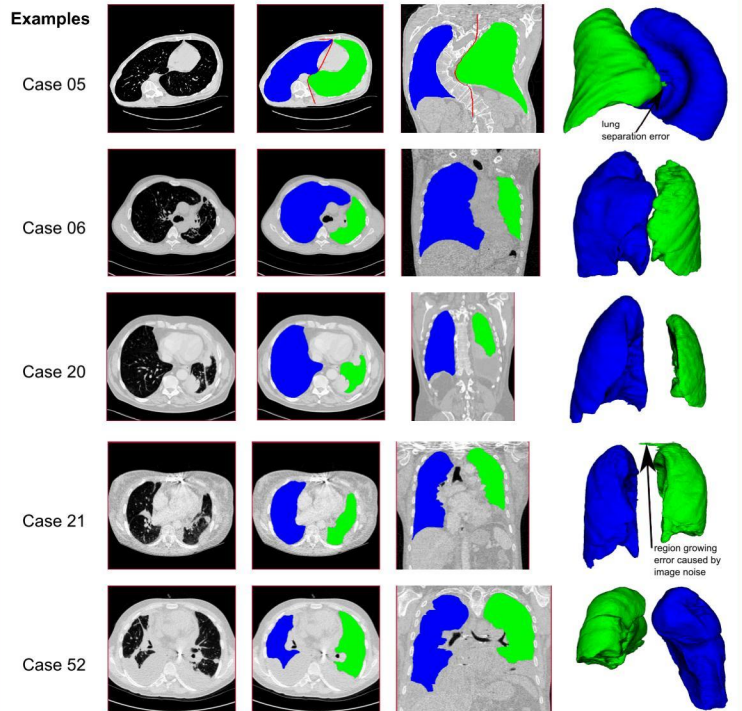
The automatic lung segmentation methodology implemented in *yacta* was applied to all 55 LOLA11-datasets. The achieved results were compared with manually generated reference segmentations. As measure the volumetric overlap fraction was used, which compares the relative volumetric overlap of two binary segmentations:  $o(V_1, V_2) = \frac{\|V_1 \cap V_2\|}{\|V_1 \cup V_2\|}$ .

The segmentation results with morphological closing (score: 0.970, step 6, see Table 1) are with respect to this measure slightly better as the results without closing (score: 0.964). The average runtime on a PC (Intel Xeon CPU, 2.83 GHz, 4GB RAM) per case was 7.3 minutes without morphological closing and 11.4 minutes with morphological closing. For details on the set-up of the study, collection of data and evaluation of segmentations see <http://www.lola11.com/>.

**Table 1.** Results of lung segmentation for the 55 scans in LOLA11 - with morphological closing.

	mean	SD	min	Q1	median	Q3	max
left lung	0.971	0.0925	0.309	0.982	0.988	0.992	0.997
right lung	0.969	0.134	0	0.986	0.99	0.993	0.998
score	0.970						

## Examples



It is worth to mention, that *yacta* is integrated (running on a virtual server) in the daily routine of the Department of Radiology at the Thoraxklinik Heidelberg. Just there, more than 2500 datasets were analysed by *yacta* until now. If errors occur in the segmentation process, they are analyzed and the algorithms are modified accordingly, difficult datasets are captured in a database. New releases of *yacta* are tested on a database containing currently more than 150 scans from different sources. This gives the possibility to develop the software step by step, without losing quality achieved already.

## Conclusion

We presented a method for fully automated segmentation of the lungs from CT scans. Our method worked well on the challenging datasets of LOLA11, nevertheless we have gathered valuable information for our future work. Generating a common database covering a wide range of possible CT scans is an important step for improving and comparing different lung segmentation methods.

Our implementation can be classified as advanced conventional method. The good score achieved with this method shows that it is a suitable solution for lung segmentation. Nevertheless there are still problems to solve and ideas to improve the method, e.g.: (1) Using a higher HU threshold for the threshold based segmentation in step 3 can lead to a better segmentation. (2) The minimal path algorithm is implemented in 2D, working on axial slices. This should be a real 3D method. (3) Integration of a robust lobe segmentation. (4) Faster runtime. (5) Better airway tracer.

## Literature

- [1] Weinheimer, O., Achenbach, T., Düber, C.: Fully automated extraction of airways from ct scans based on self-adapting region growing. In: Proceedings of the Second International Workshop on Pulmonary Image Analysis. (2009) 315–321
- [2] Mueller, D.: Fast marching minimal path extraction in itk. The Insight Journal (2008)

## Further Information

Please contact [mail@oliwe.com](mailto:mail@oliwe.com). More information on this and related projects can be obtained at [www.oliwe.com](http://www.oliwe.com). You can find an online PDF-version of this poster here: [www.oliwe.com/publications/Weinheimer\\_2011\\_LOLA11\\_Poster.pdf](http://www.oliwe.com/publications/Weinheimer_2011_LOLA11_Poster.pdf)

