# COMPUTER AIDED DIAGNOSIS LUNA16: CANDIDATE DETECTION

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### **ABSTRACT**

Write your abstract here.

### 1. INTRODUCTION

In the *LUng Nodule Analysis 2016 (LUNA16)* challenge, we aim to detect lung nodules in low-dose lung CT images. Firstly, we have segmented the lungs from the images. The next step is identifying candidates for lung nodules. Our aim is to detect enough candidates to include all actual lung nodules. In other words, be as sensitive as possible. The future step is of course to remove the false positives, but first we strive towards including the annotated nodules.

We continued working on the deep learning approach we used for the previous phase. In that phase we worked on lung segmentation and for this phase we treated candidate detection again as a segmentation problem. Secondly, we also used an image processing approach where we use the features of nodules such as blobness for candidate detection. These approaches and results are explained more detailed in the following sections.

Table 1. Caption here.texttexttexttexttexttext

You can include figures as follows:

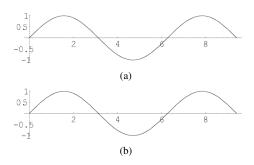


Fig. 1. Caption here.

### 2. METHOD

Explain the proposed method here. You can use subsections to detail each part of your approach as follows.

### 2.1. Fully convolutional network

Your text here.

### 2.2. Nodule detection using blobness feature

We used a second approach for detection lung nodule candidates which is the more conventional approach of using their blob-like structure. The first step is applying the lung segmentation as a mask on the lung CT images to reduce the search space to the actual lungs. The algorithm for the actual candidate detection is as follows:

- Apply two different sized Laplacian of Gaussian (LoG) filters. A LoG filter is circular-symmetric and thus widely used for blob detection, but has a maximum response to a particular blob size. We combined two filters with normalized kernels (IS THIS NORMALIZED?), the first with  $\sigma=3$  and  $kernel\ extent=7$  and the second with  $\sigma=5$  and  $kernel\ extent=11$  CHECK VALUES. We chose these parameters based on the paper of Van de Leemput et al. [?]
- Take the pixel-wise maximum of the results of the LoGs to combine them to one result.
- Threshold the resulting image with value THRESH-OLDVALUEAFTERLOGS. See Figure 2 for an example result. This value is established by sampling the values of known nodules and healthy lung tissue.

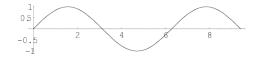
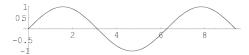


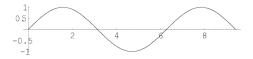
Fig. 2. The thresholded image after taking the maximum of the two LoGs.

 Do a connected component analysis on the resulting image, and select only components with values between CONNECTEDCOMPONENTSVOLUMEVAL-UES. This identifies the initial candidates. See Figure 3 for an example result.



**Fig. 3**. The result of applying a connected component analysis on the image shown in Figure 2.

- Threshold the resulting image again with value THRESH-OLD2VALUE.
- Apply a Hough transform to identify unwanted candidates, i.e. candidates with features not corresponding to lung nodule characteristics.
- Threshold the resulting image again to remove the candidates identified by the Hough transform and keep the lung nodule candidates. See Figure 4 for an example result.



**Fig. 4**. An example result after applying the Hough transform and thresholding.

**ADD IMAGES** 

## 3. RESULTS

### 3.1. Conventional method

The results are based on the first thirty images in subset 9 of the dataset. The results are evaluated in two ways: the dice score of the predictions to the ground truth candidates given for this phase of the assignment, and the recall between the same datasets. The reason that the results are only examines on thirty images is that this method is very time consuming (over 10 minutes per image) and the method was completed relatively close to the deadline.

The average dice score obtained by this method is ME-ANDICESCORE, with a standard deviation of STDDICESCORE. This score is relatively low, compared to score reaches in the previous phase, but the main goal of this method is to find all posible nodules. Thus, a large recall value is more important than a large dice score.

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The recall obtained with this method is RECALL.

3.2. fcn

### 4. DISCUSSION

### A. CONTRIBUTIONS

Luc Nies:

Tom van de Poll:

**Harmen Prins:** 

**Steven Reitsma:** 

**Inez Wijnands:** Contributed to creating the MeVisLab candidate detection method using blob detection. Wrote introduction section and method section on the conventional approach in report.

### **B. REFERENCES**

[1] S. van de Leemput, F. Dorssers & B.E. Bejnordi (2015). A novel spherical shell filter for reducing false positives in automatic detection of pulmonary nodules in thoracic CT scans. *Proceedings SPIE Medical Imaging*, 9414:94142P.