

ISMI – NODE21 Challenge

Nodule Generation Track

Group 5

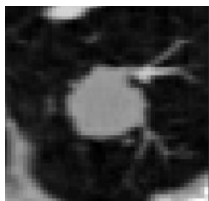
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Generation synthetic nodule

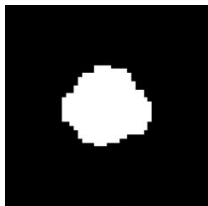
- The aim of the task is producing more nodules to enhance the dataset.
 - This could improve performance of detection systems.
 - This could help detecting cancer in earlier stages.
-
- **RQ:** Is using CT patches feasible to generate X-Ray nodules?

Materials

- Applying different sources of data:
 - 1186 CT patches containing nodules and nodule masks.
 - 1031 frontal X-Ray images and lung masks¹.



An example of CT patch



An example of Mask

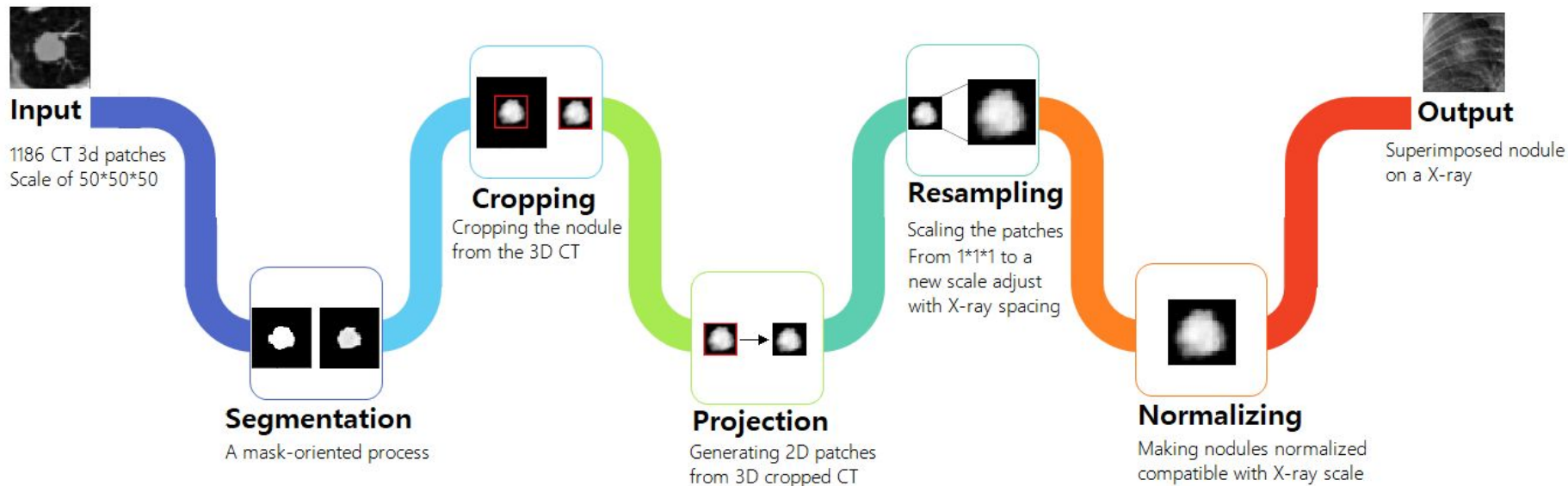


An example of X-Ray

Method

- Creating a pipeline in Python mostly inspired by Litjens et al. work
 - Feeding the pipeline with 1186 CT in scale of 50*50*50
 - All CT voxels are **segmented** using the associated nodule mask.
 - The segmented CT is **cropped** to have 3D nodule, before projecting.
 - The 3D cropped nodule is **projected** to a 2D image (Campo et al.).
 - The 2D nodule is **resampled** using interpolation to an isotropic resolution equal to the X-ray.
 - Generated nodule is **augmented** by four different techniques to increase the training data.
 - The pixel value of nodules is **normalized** to get compatible with X-ray range.
 - The generated nodule is **superimposed** onto the X-ray image (randomly chosen location)

Preprocessing



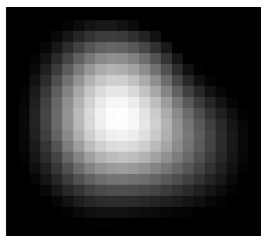
Superimposing Phase

- We applied an empirical mathematical method.
- The method named the Perfect Spherical Nodule Simulation.
- Weighting central pixels of nodules more than borders.
- Methods parameters guarantee proper blending especially near the edges.
- Each pixel value of nodule, after multiplying by c , would add with X-ray corresponding pixel.

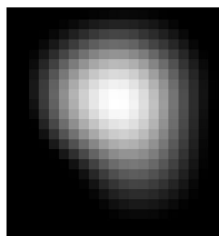
$$c[r] = C(\frac{4}{D^4}r^4 - \frac{4.2}{D^2}r^2 + 1)$$

Data Augmentation

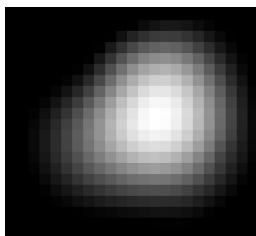
- Since new synthetic data needs to be practical, making the options of augmentation methods are limited.
- We applied four variant augmenting methods
- Rotation, Vertical and Horizontal Flipping, and Zooming-in
- Generated 930 nodules



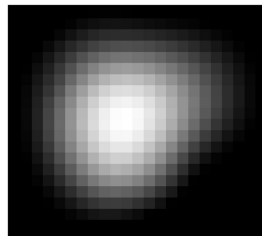
(a) Original Version



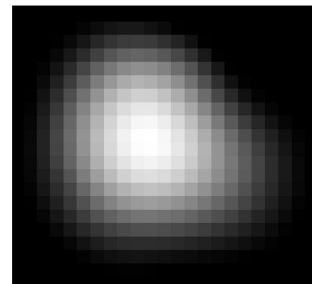
(b) Rotation 90



(c) Flip Horizontally

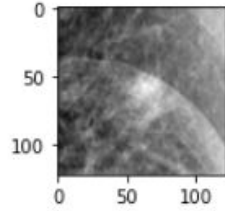
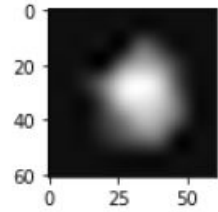


(d) Flip Vertically

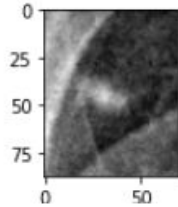
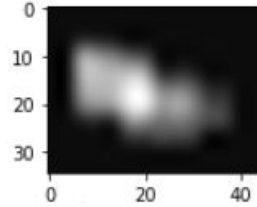


(e) Zoom-in

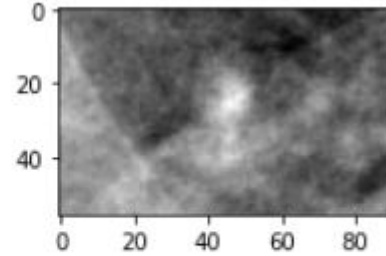
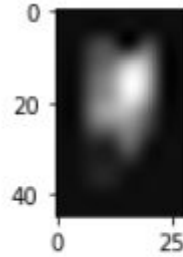
Generated and Superimposed Nodules



(a)



(b)

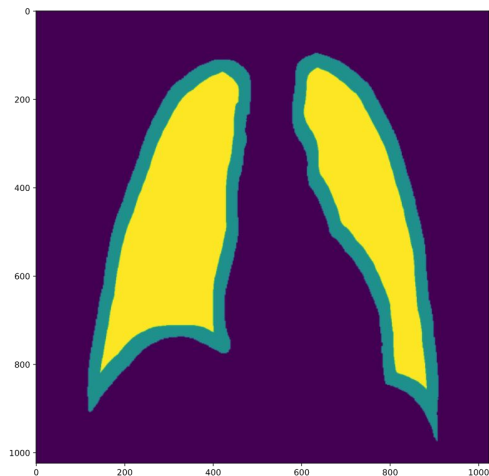


(c)

Two examples of projected nodules and superimposing nodules

Lung Mask Preprocessing

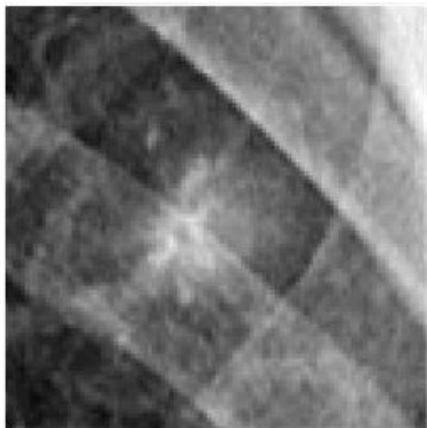
- A specific location of X-ray must be chosen to superimpose the 2D nodules.
- The numpy random number generator was applied.
- To prevent inserting nodule outside of lungs, lung mask was used to crop them.



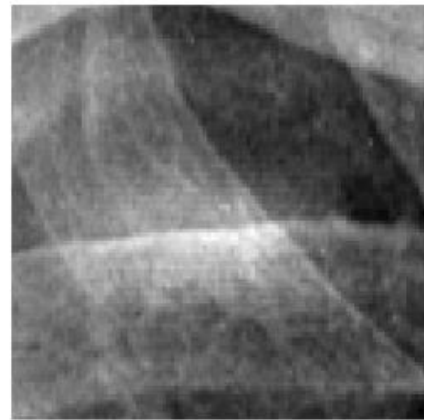
Final Result



(a)



(b)



Two examples of superimposed X-rays

Detection Phase

- Detection is necessary to test the performance enhancing capabilities of our newly generated nodule dataset.
- We tested the data on an already existing deep learning detection method.
- We chose Faster RCNN that is quick to implement.
 1. Training the model on only the real data, the outcome of this would serve as a baseline performance.
 2. Training the model on only generated data.
 3. Training the model on dataset which was 50 percent real data and 50 percent generated data.

The Experimental Setup

- Superimposing the projected nodules in different sizes
- ??? iteration to superimpose nodules in 1031? CXR.
- Testing the augmented results with a Faster RCNN? detection algorithm.
- Splitting dataset to training, validation, and test set (90-5-5 %)
- We used a learning rate of 0.005, momentum of 0.9 and a weight decay of 0.0005.
- We trained the model in 10 epochs.

Result

- Each of the systems was tested on 52 x-rays of real patient data.
- the mAP* of the models to detect nodules was calculated.

Dataset	mAP
Real	0.101
Generated	0.0
Mixed	0.039

* mean Average Precision

Discussion

- We could insert different size of nodules.
- We generated much more data than the first time
- We modified the resampling based on the meaning of pixel size (in CT and X-ray)
- We opted a superimposed model in which we added nodule into X-ray instead of computing the mean.
- We modified the normalization process
- We added data augmentation
- We could train a model with our generated data

Future Works

- Choosing a more correlated location on X-ray to inset nodules (it is already randomly chosen)
- Generating synthetic nodule to add behind the heart or diaphragm
- Enhancing the number of data by merging data augmentation techniques
- Applying Gans to produce nodule