

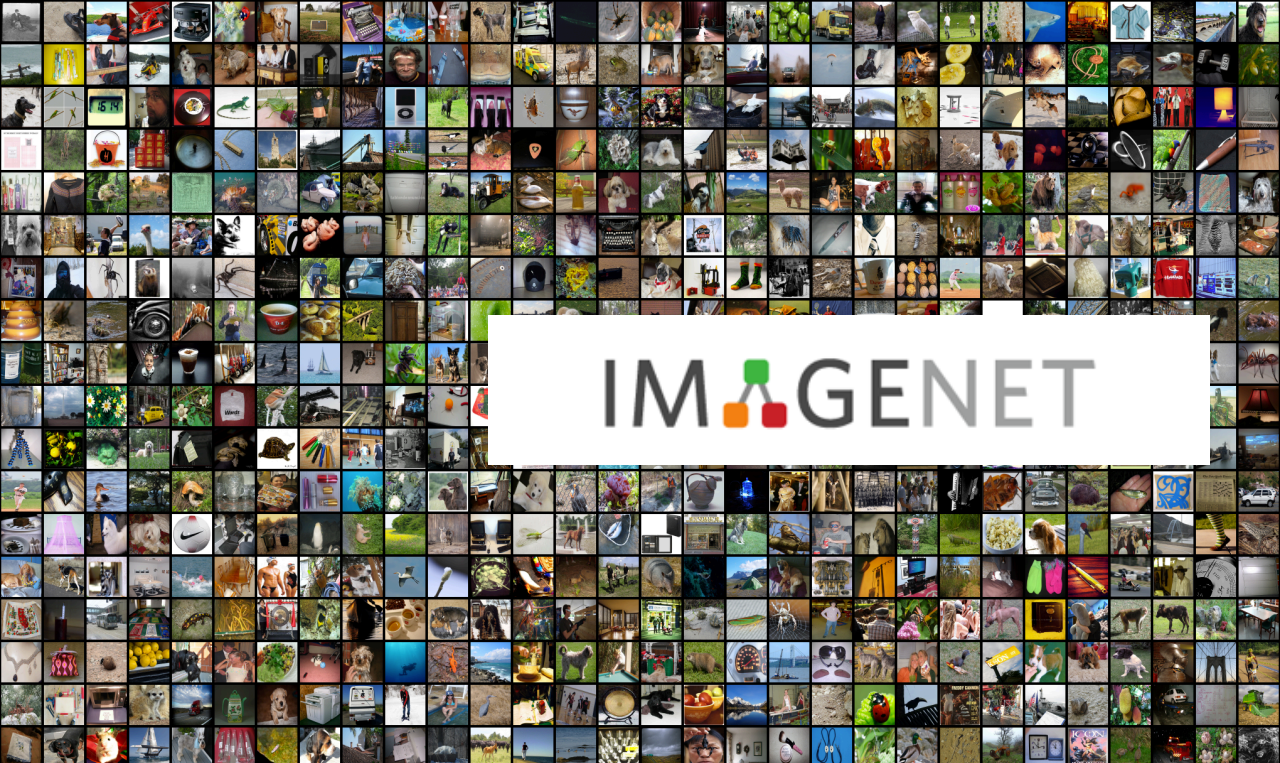
We provide pre-trained 3D models!

Models Genesis: Generic Autodidactic Models for 3D Medical Image Analysis

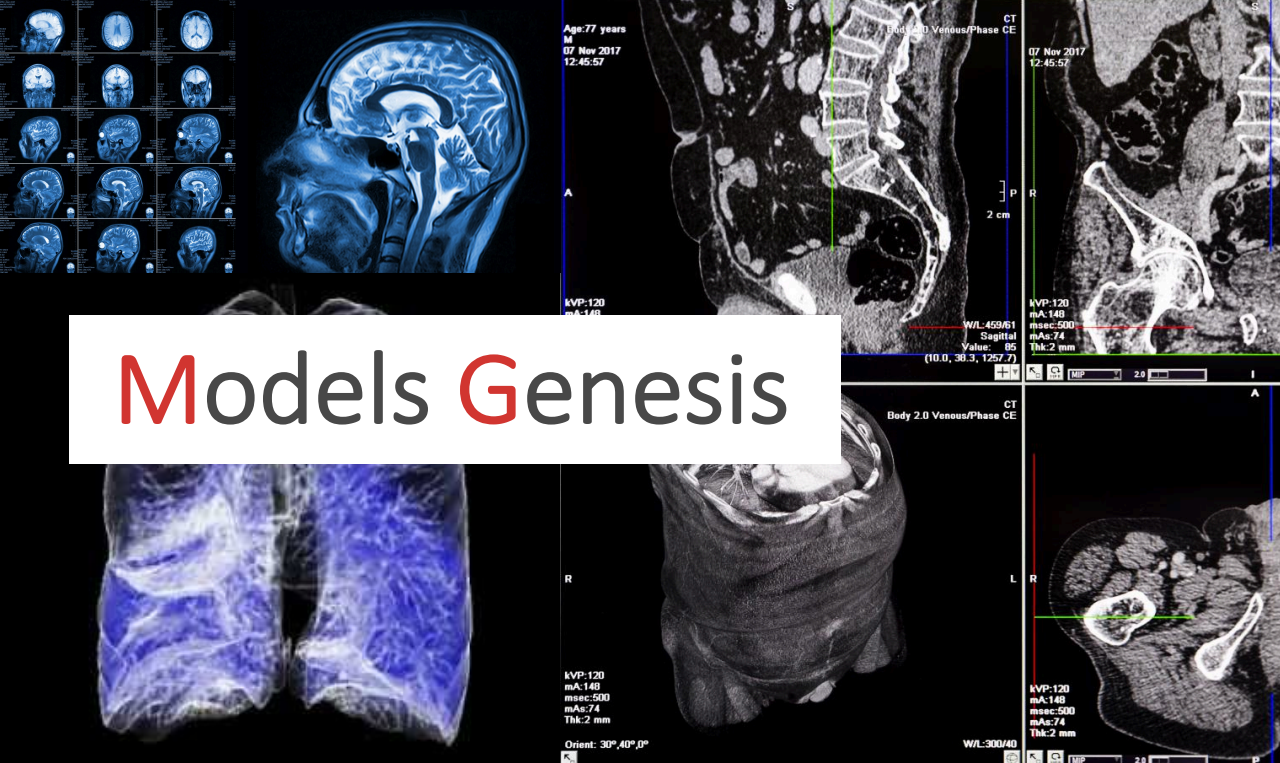
Zongwei Zhou¹, Vatsal Sodha¹, Md Mahfuzur Rahman Siddiquee¹,
Ruibin Feng¹, Nima Tajbakhsh¹, Michael B. Gotway², and Jianming Liang¹

¹ Arizona State University

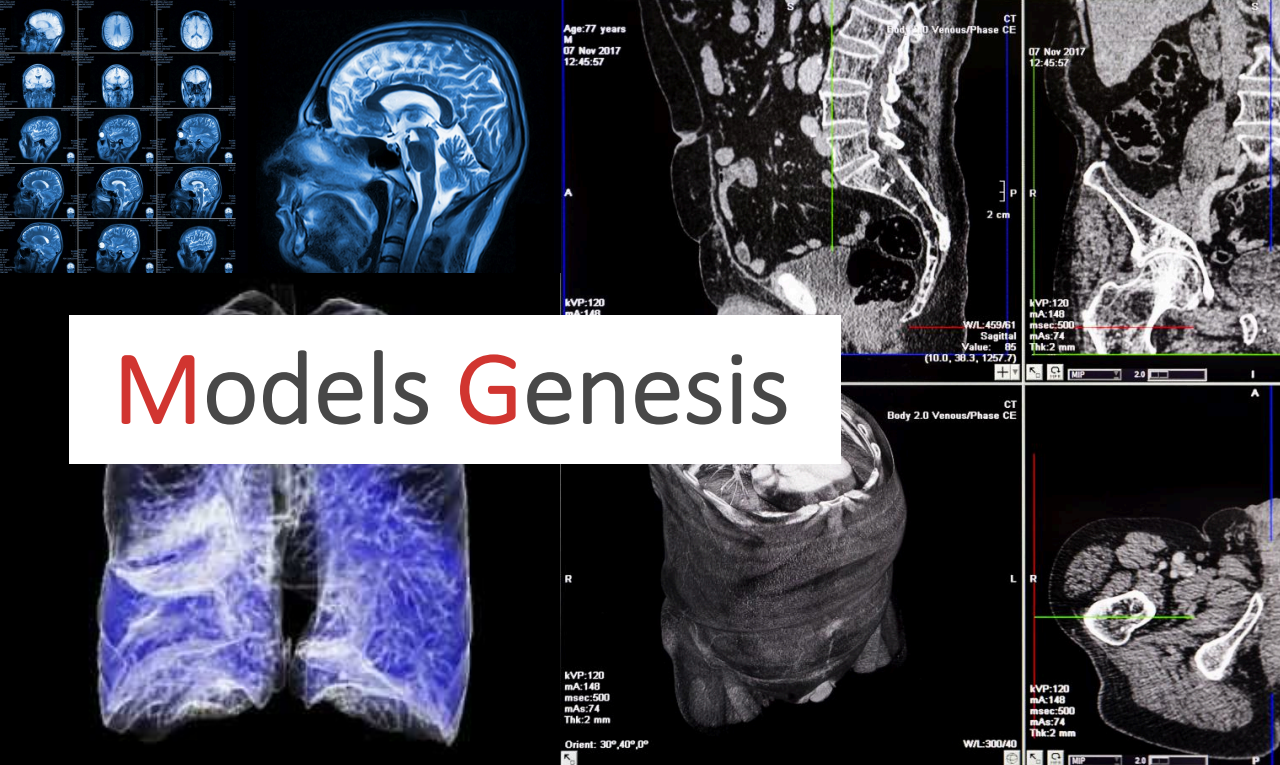
² Mayo Clinic



IMAGENET



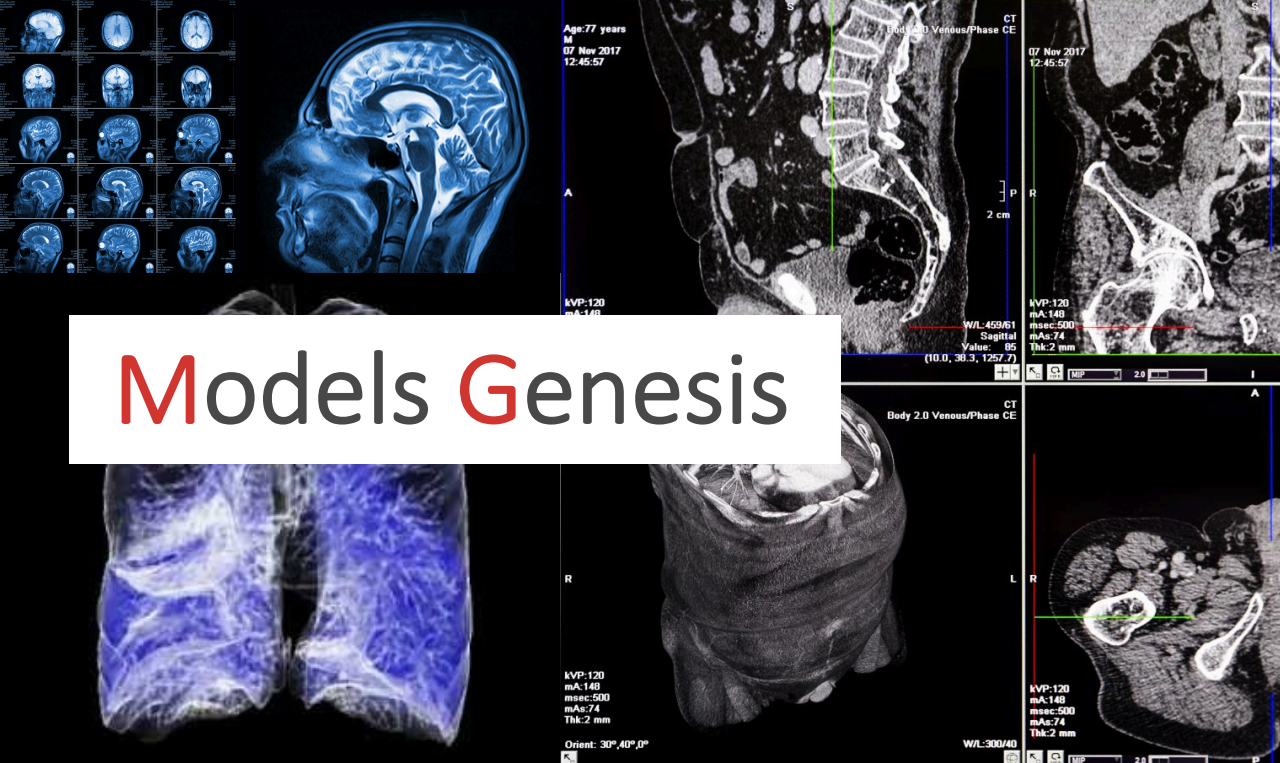
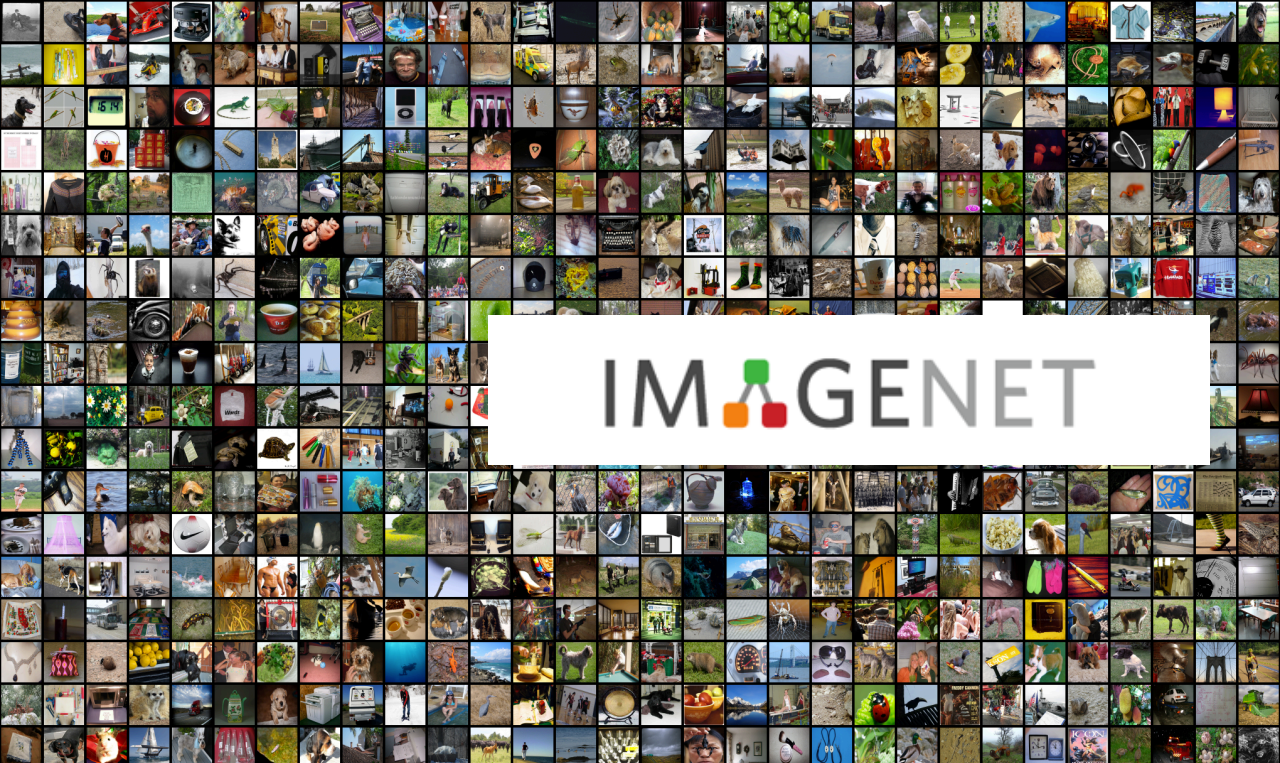
Models Genesis



Natural images

Medical images

Transfer learning: medical images → medical images > natural images → medical images



IMAGENET

Models Genesis

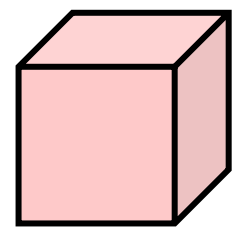
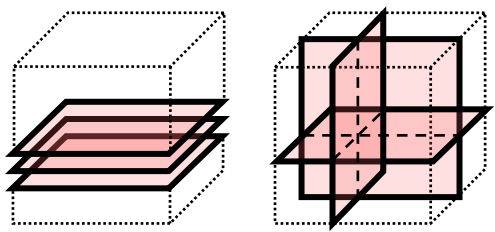
Natural images

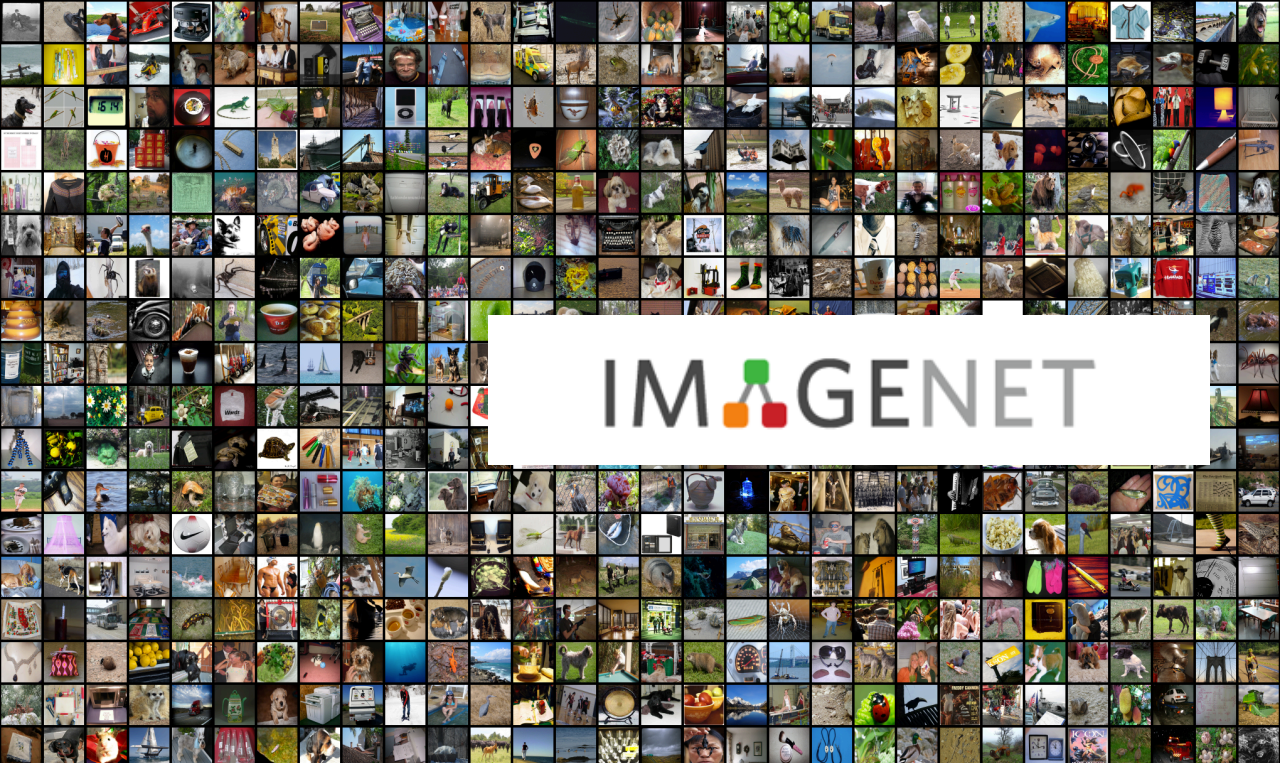
Medical images

Formed in 2D

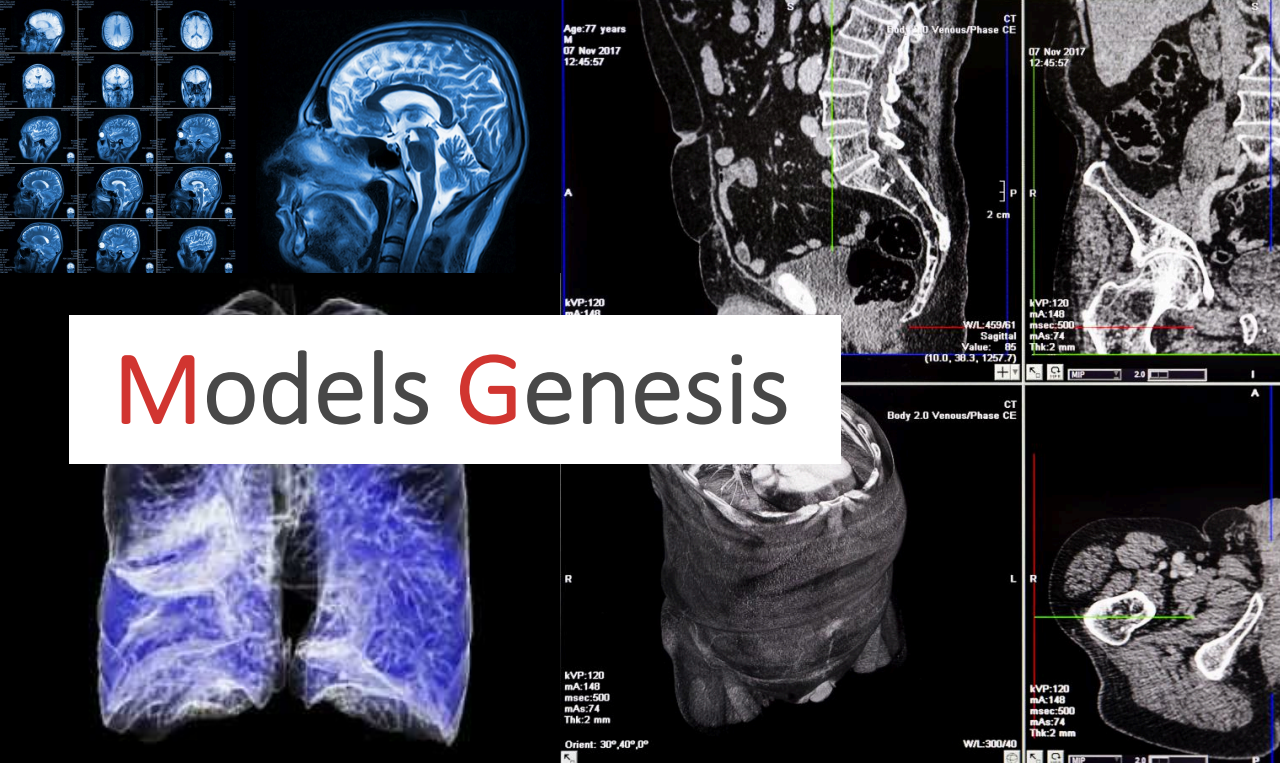
Formed in 3D

3D imaging tasks should be solved in 3D





IMAGENET



Models Genesis

Natural images

Medical images

Formed in 2D

Formed in 3D

>14,000,000 annotation

Zero annotation

Annotating biomedical images is time consuming and demanding of costly, specialty-oriented knowledge

Image Restoration Task

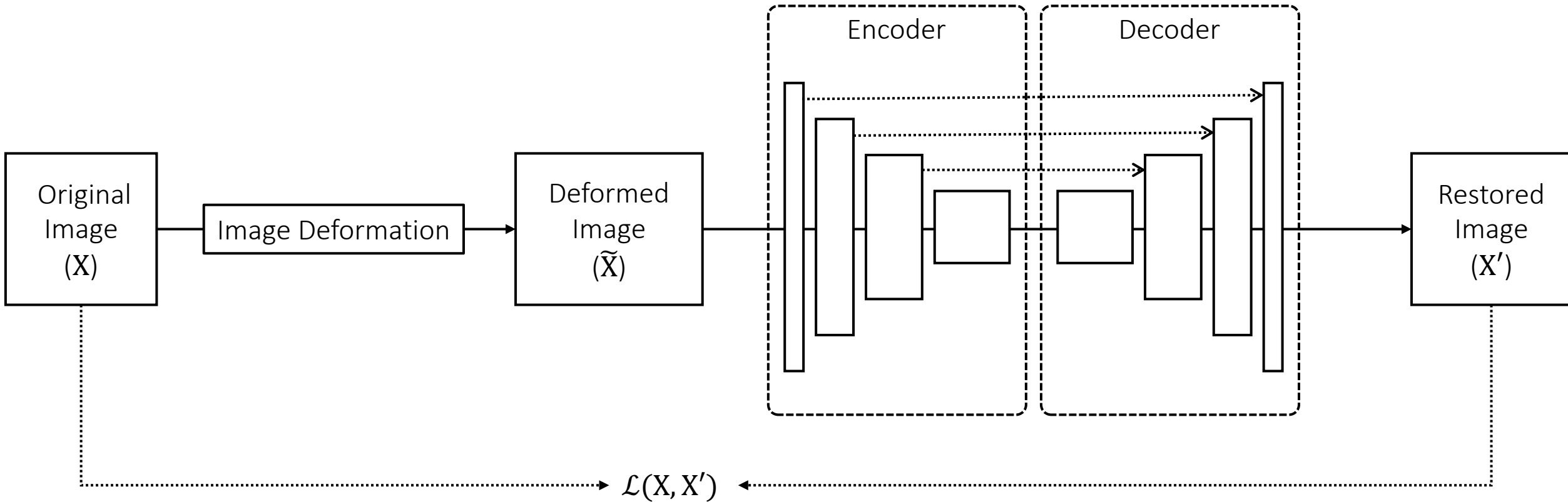
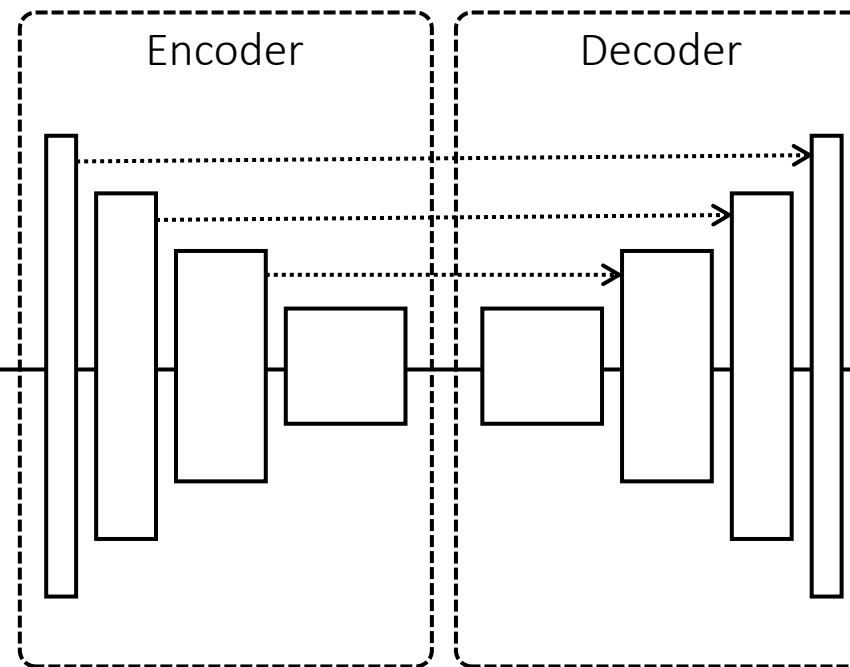




Image Restoration Task



Original Image
(X)

Image Deformation

Deformed Image
(\tilde{X})

Encoder

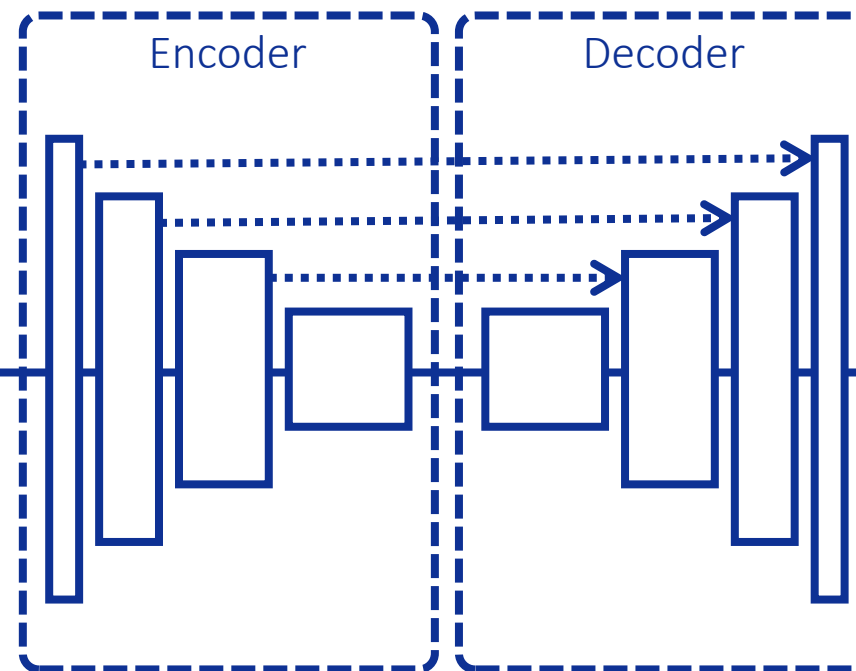
Decoder

Restored Image
(X')

$\mathcal{L}(X, X')$



Image Restoration Task



Original Image (X)

Image Deformation

Deformed Image (\tilde{X})

Encoder

Decoder

Restored Image (X')

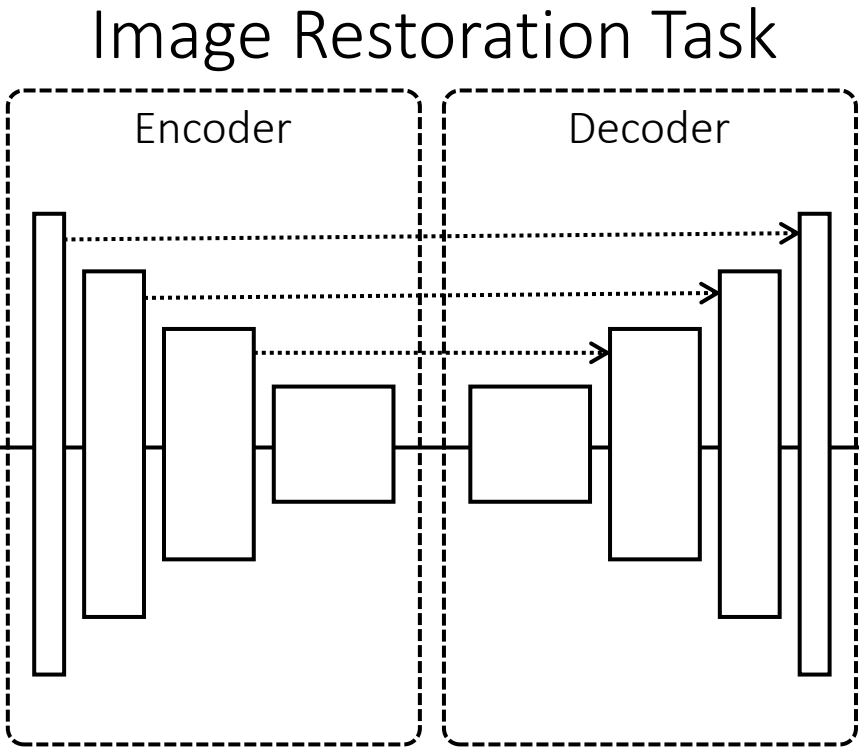
$$\mathcal{L}(X, X')$$

- → Non-linear
- → Local shuffling
- → Out-painting
- → In-painting
- → More ...

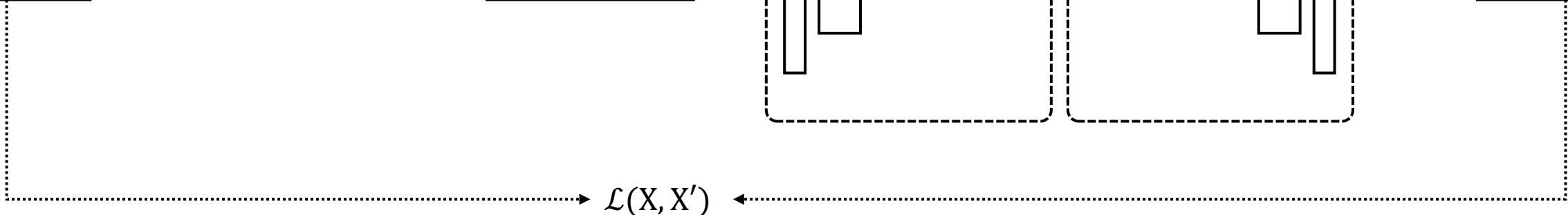
Original Image (X)

Image Deformation

Deformed Image (\tilde{X})



Restored Image (X')



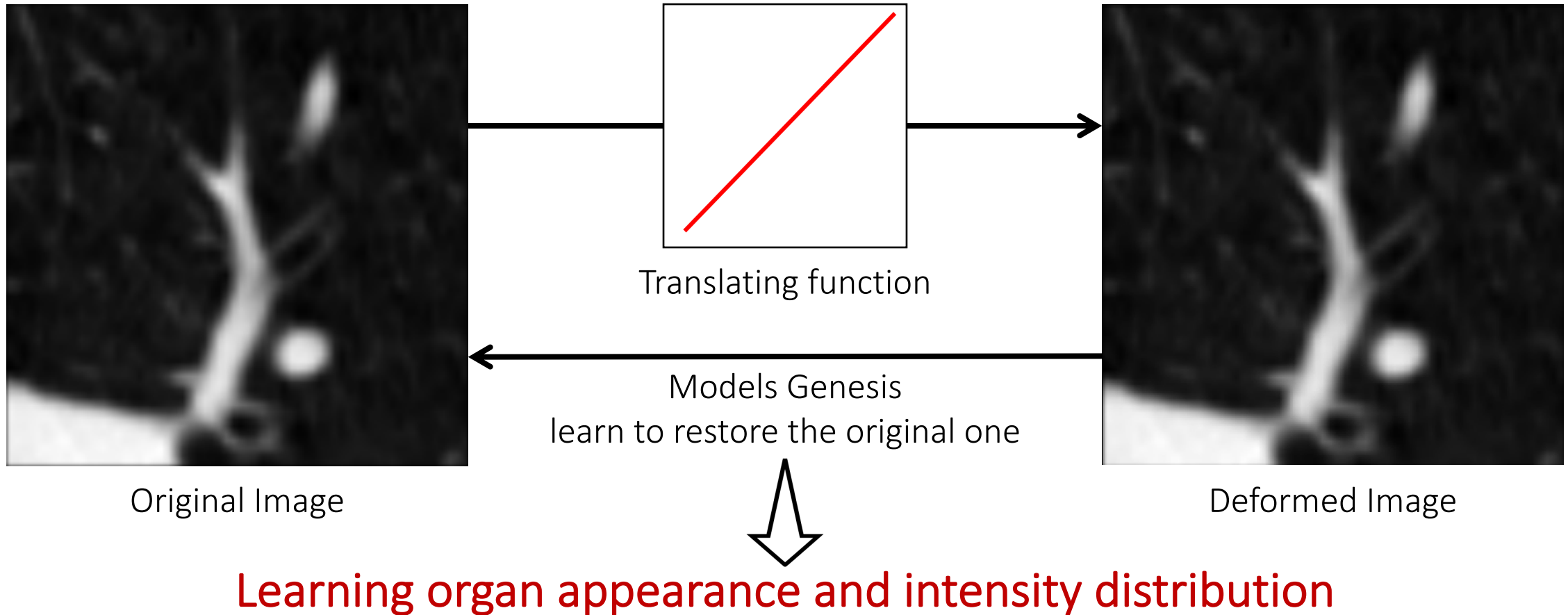
CT scan itself naturally comes with
the *pixel-wise* annotation

I. Non-linear transformation

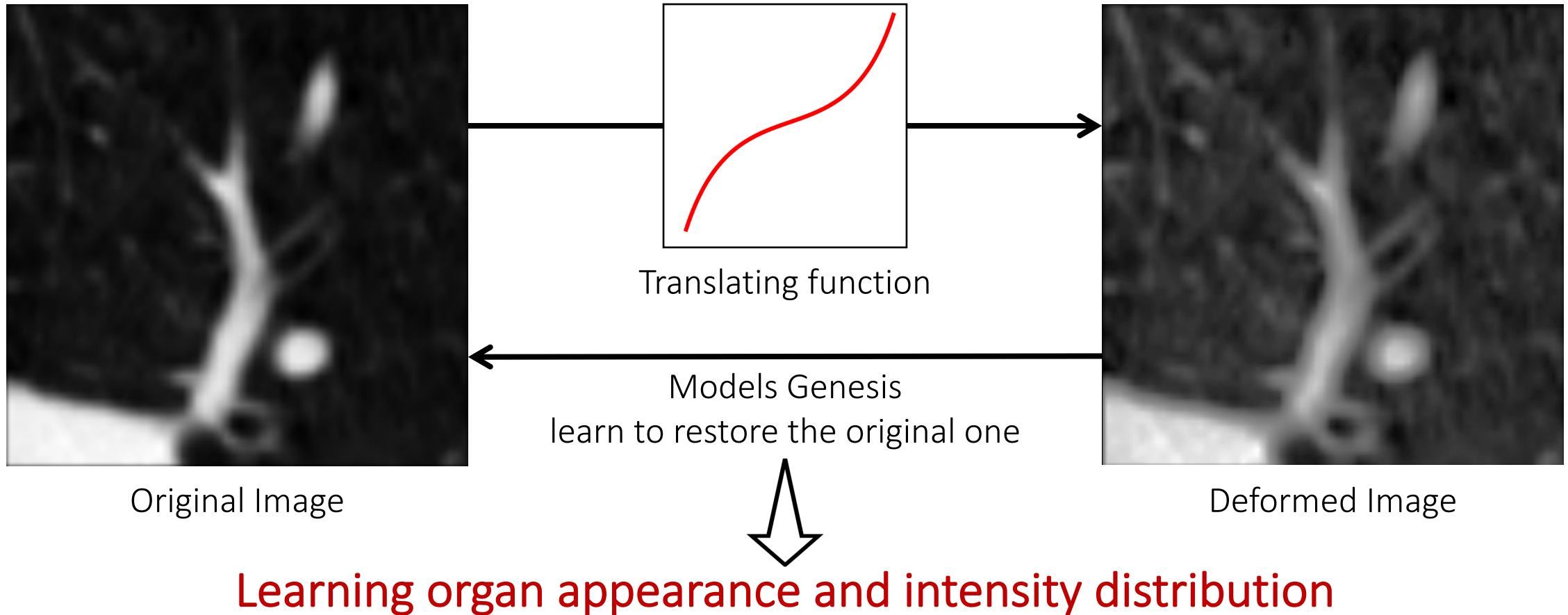
| Substance | | Hounsfield units (HU) |
|------------|-------------|-----------------------|
| Air | | -1000 |
| Fat | | -120 to -90 |
| Water | | 0 |
| Bone | Cancellous | +300 to +400 |
| | Cortical | +1800 to +1900 |
| Parenchyma | Lung | -700 to -600 |
| | Kidney | +20 to +45 |
| | Liver | +54 to +66 |
| | Lymph nodes | +10 to +20 |
| | Muscle | +35 to +55 |

Source from en.wikipedia.org/wiki/Hounsfield_scale

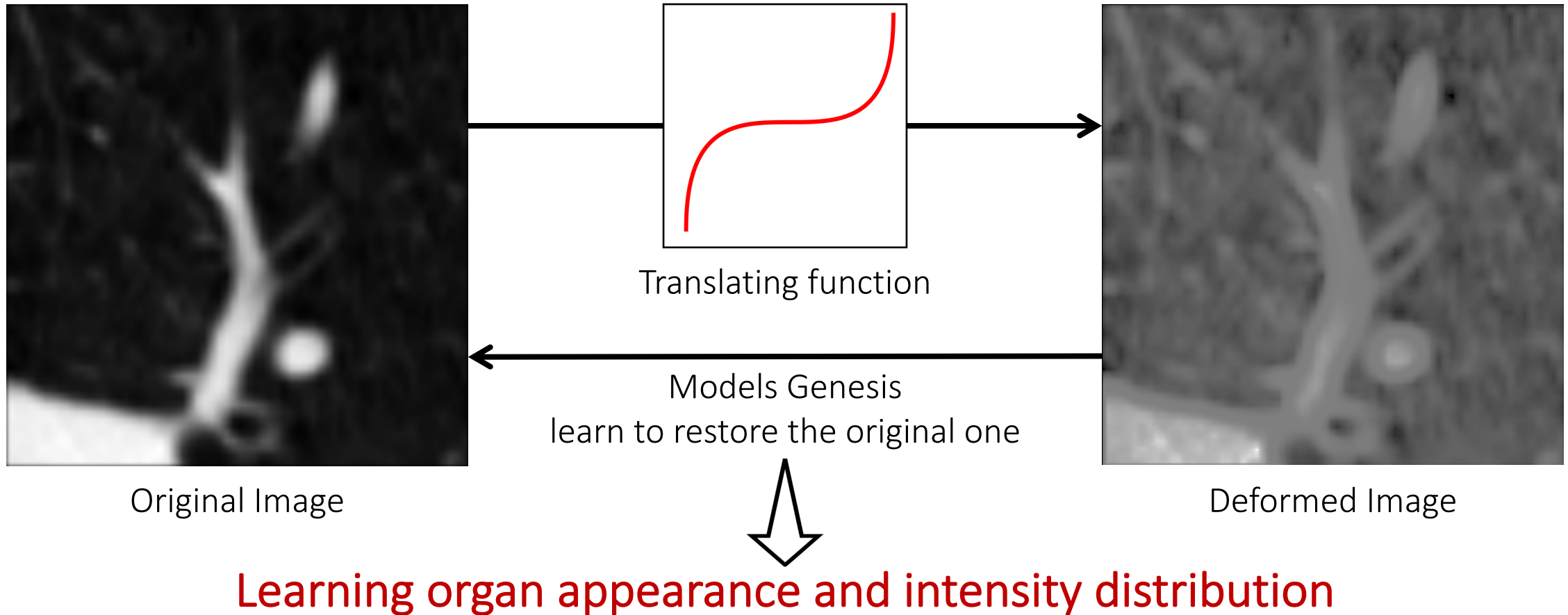
I. Non-linear transformation



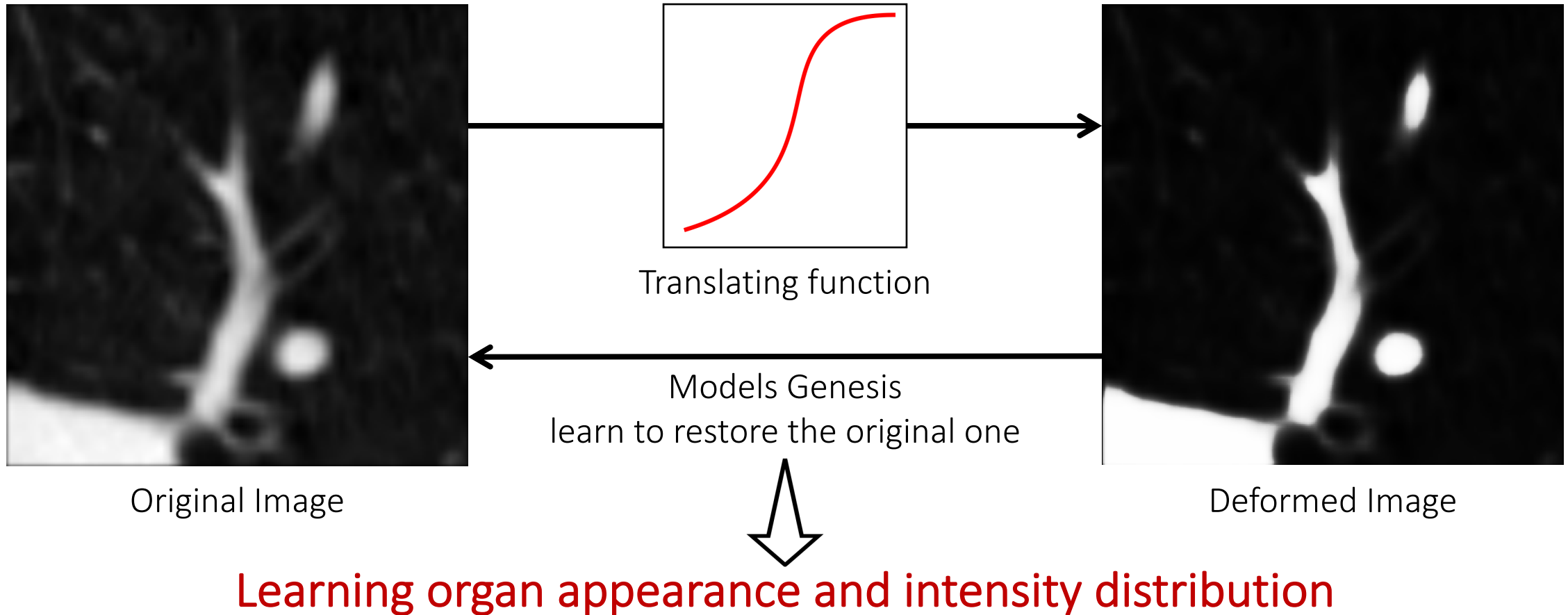
I. Non-linear transformation



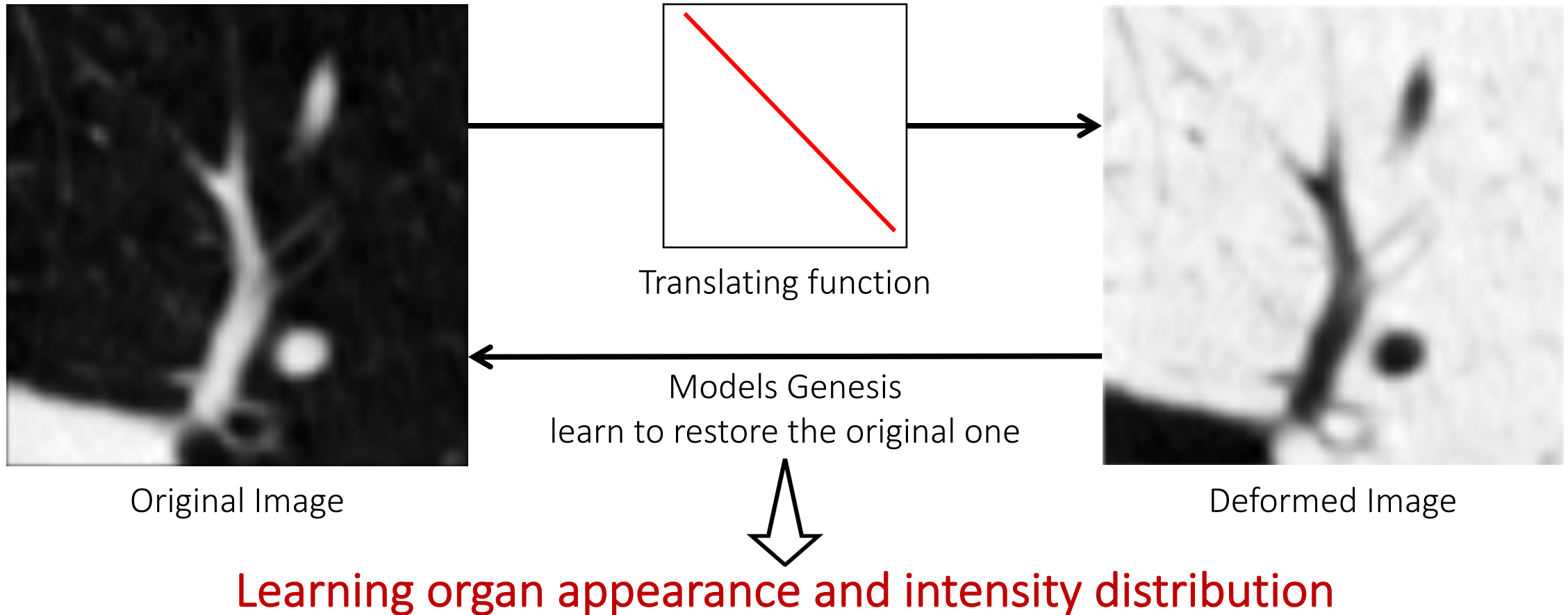
I. Non-linear transformation



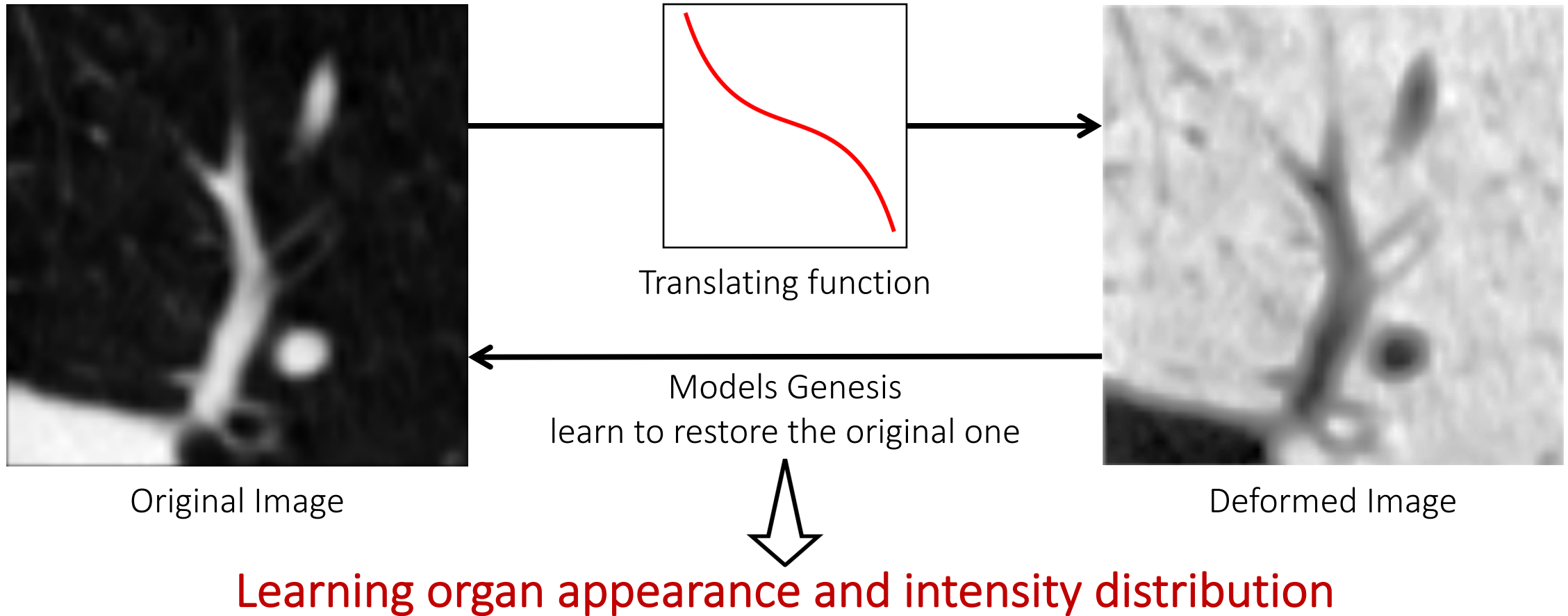
I. Non-linear transformation



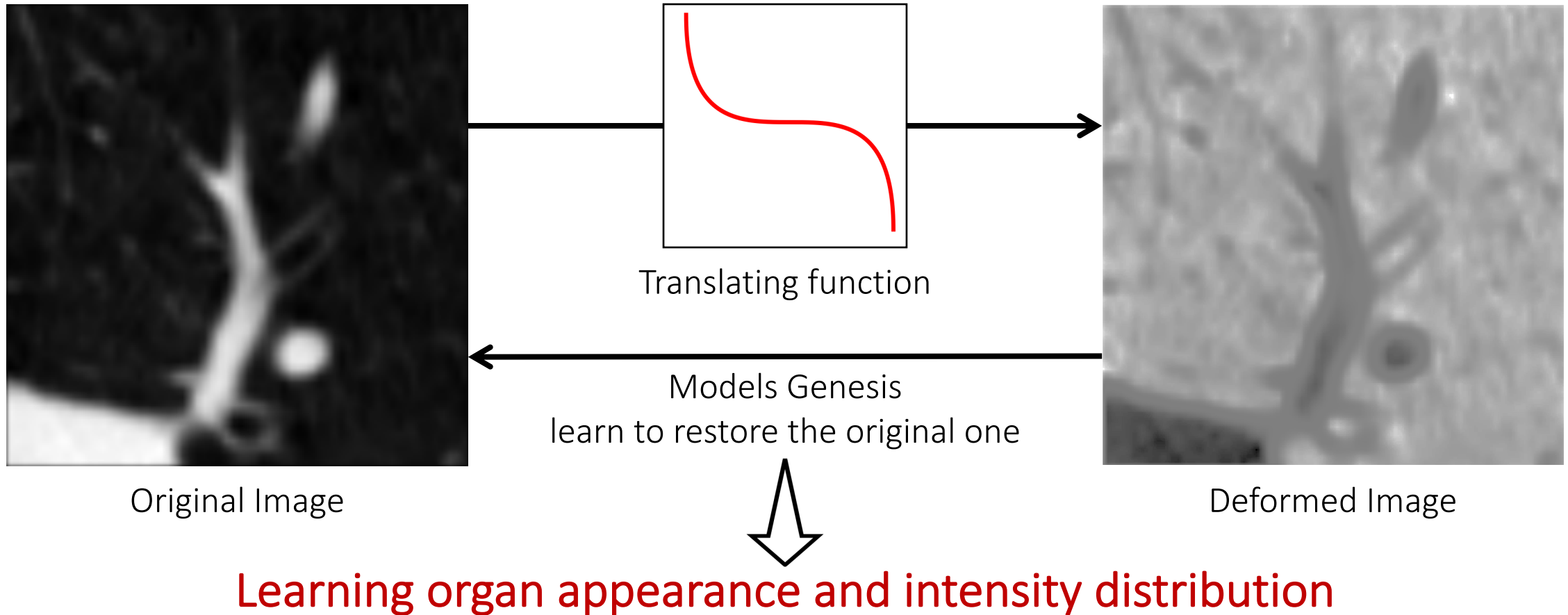
I. Non-linear transformation



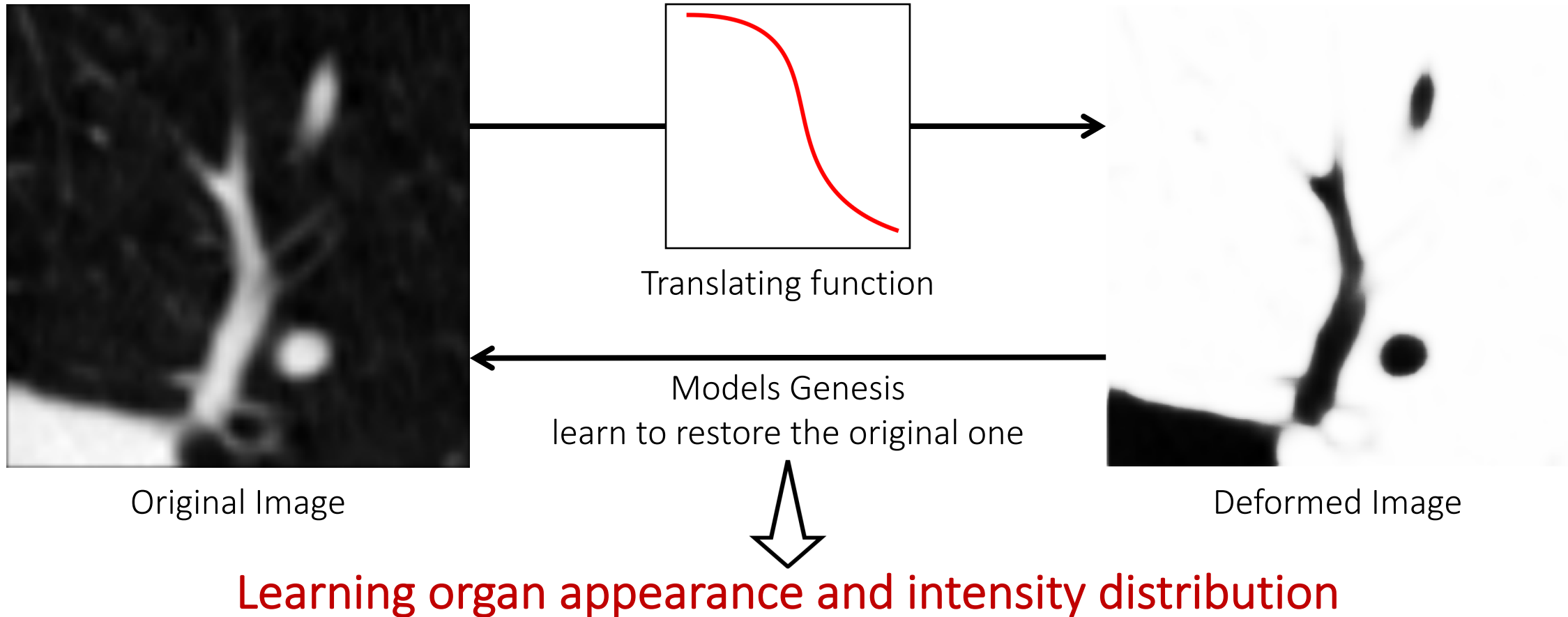
I. Non-linear transformation



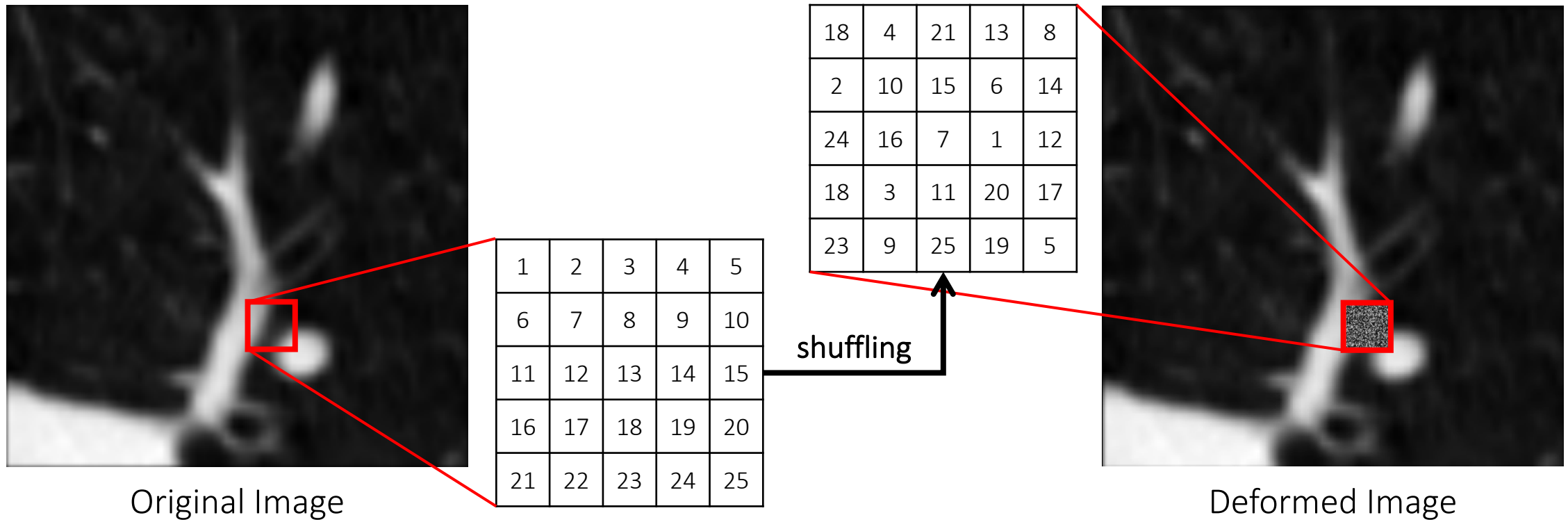
I. Non-linear transformation



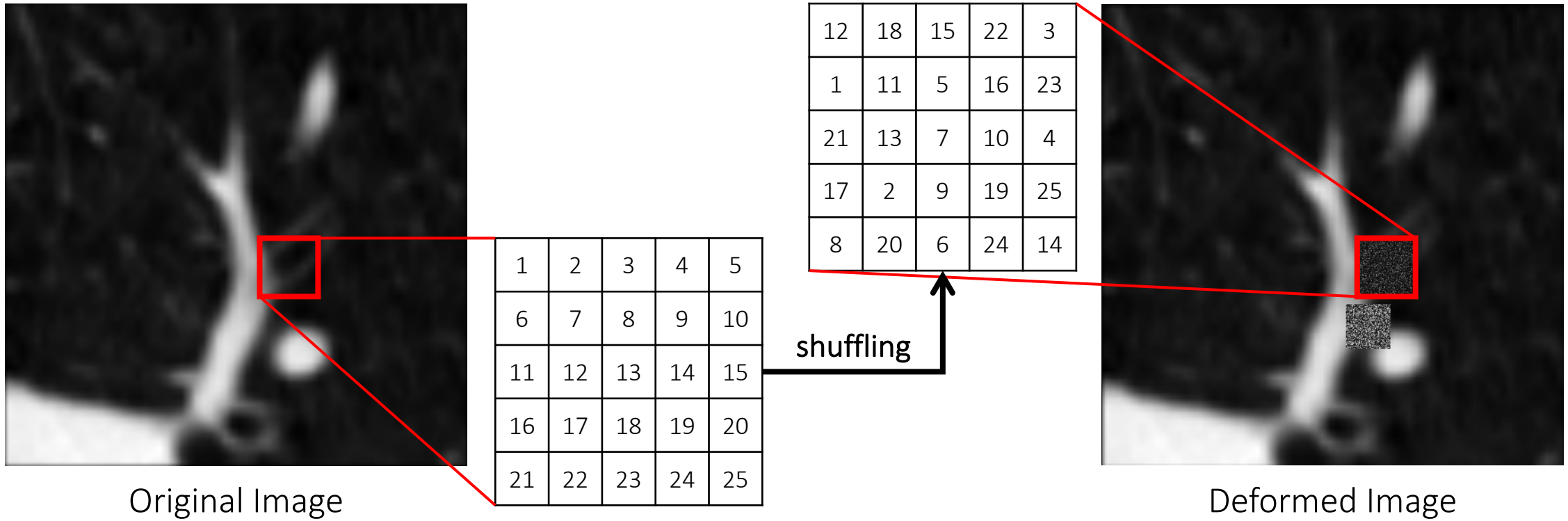
I. Non-linear transformation



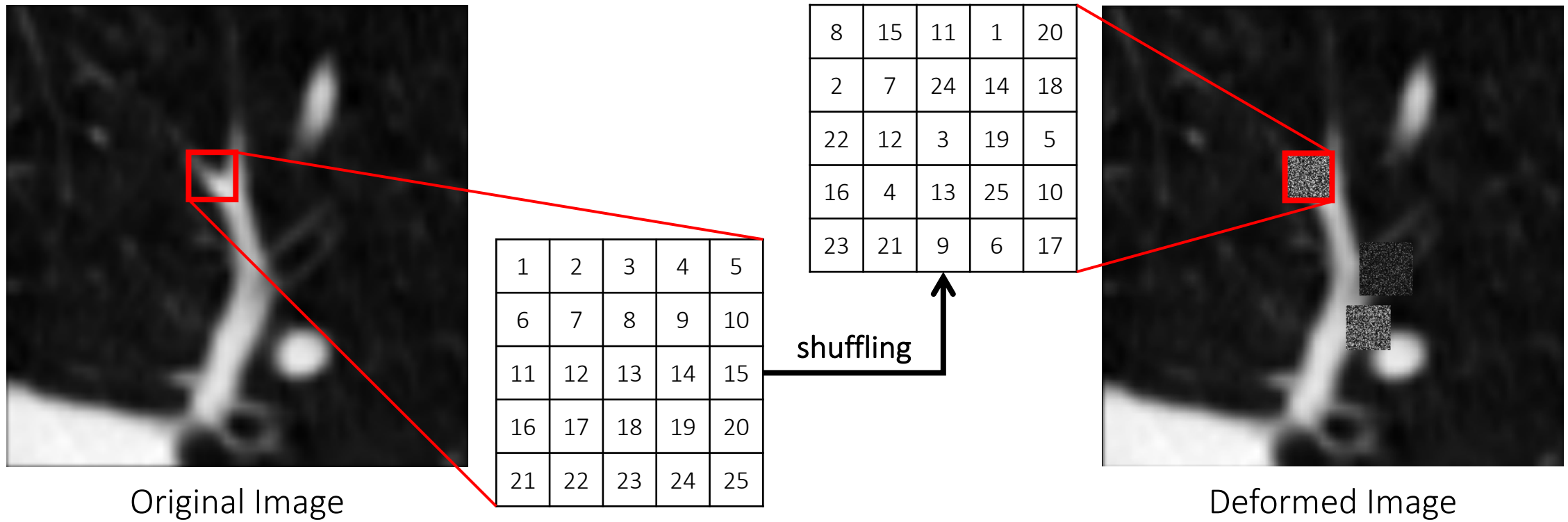
II. Local pixel shuffling



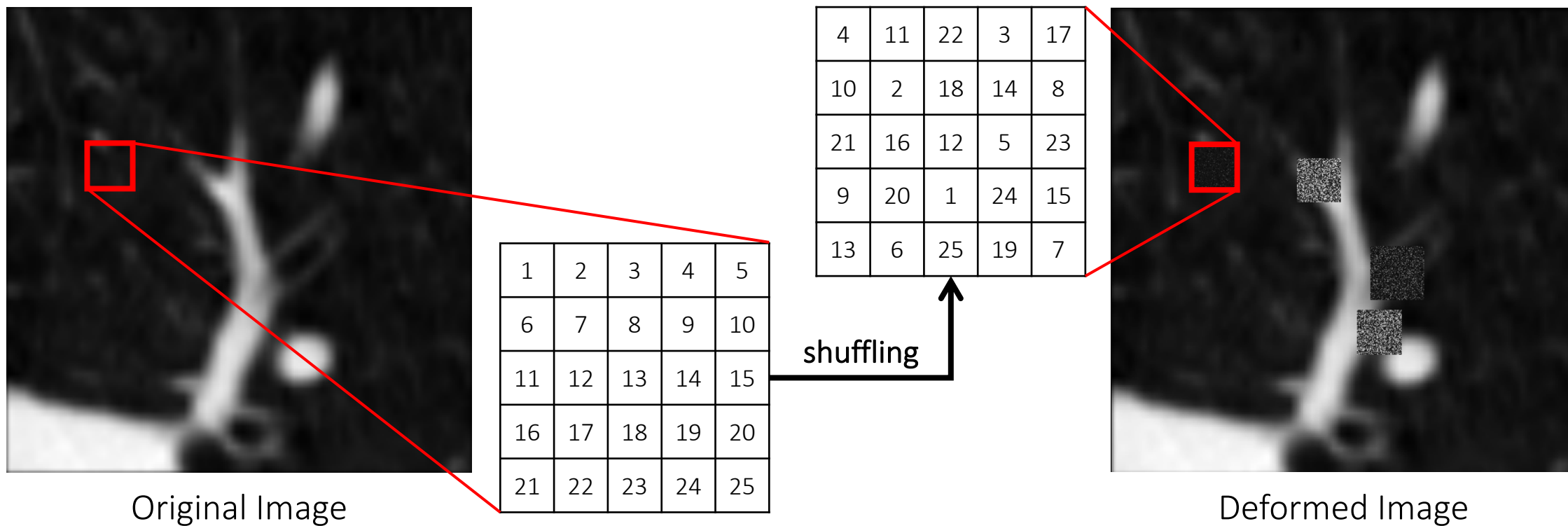
II. Local pixel shuffling



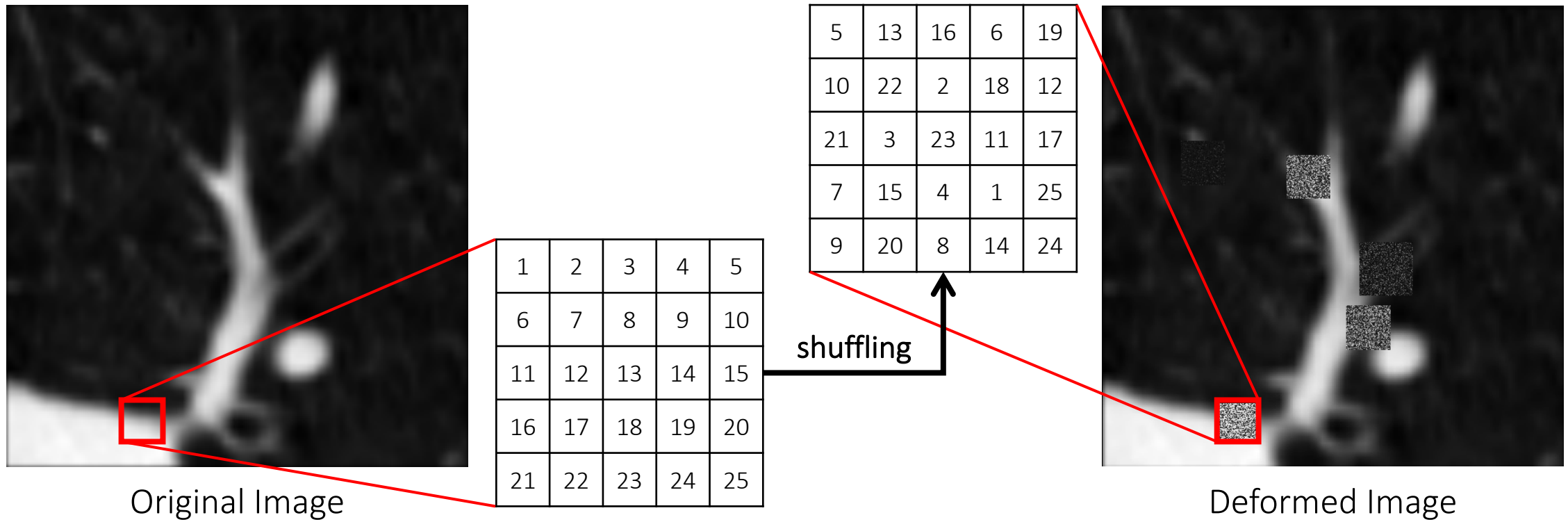
II. Local pixel shuffling



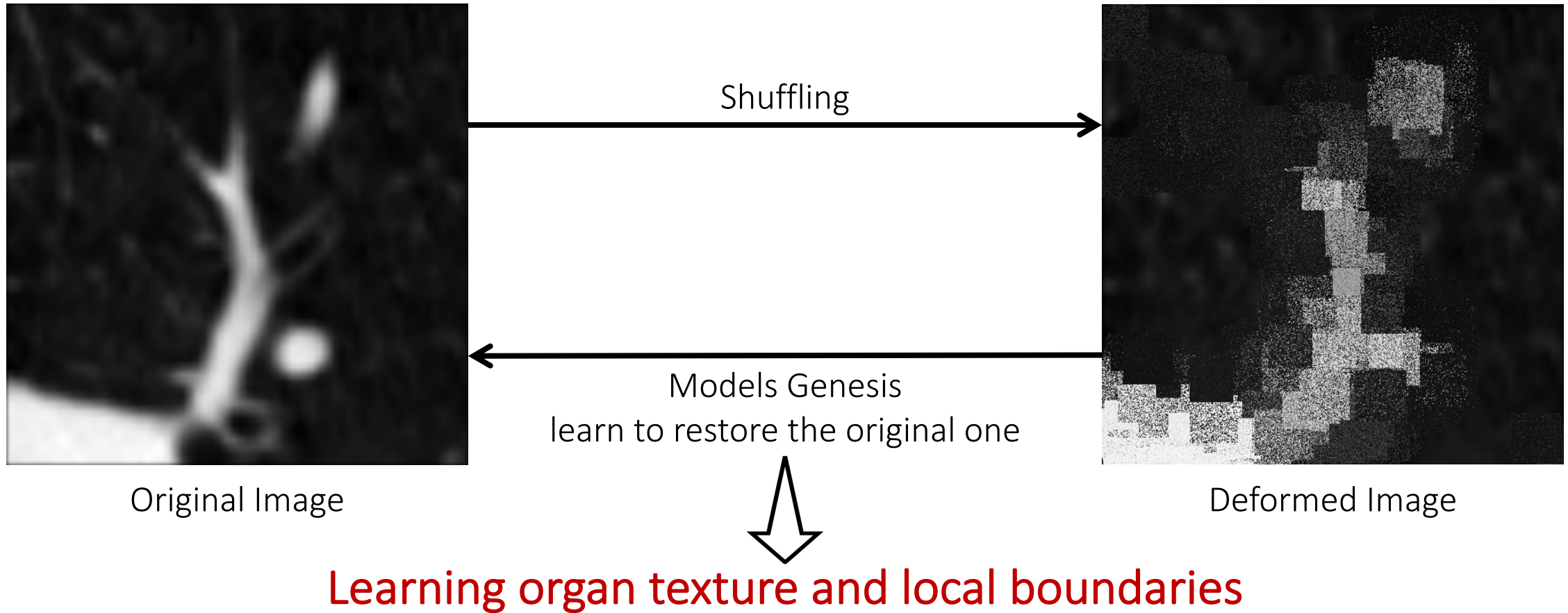
II. Local pixel shuffling



II. Local pixel shuffling

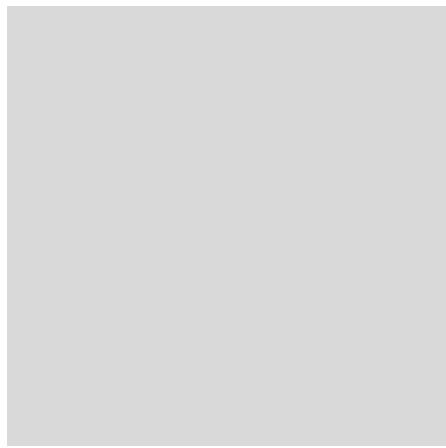


II. Local pixel shuffling

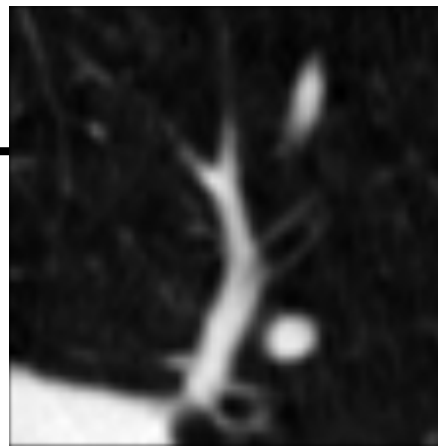


III. Out-painting

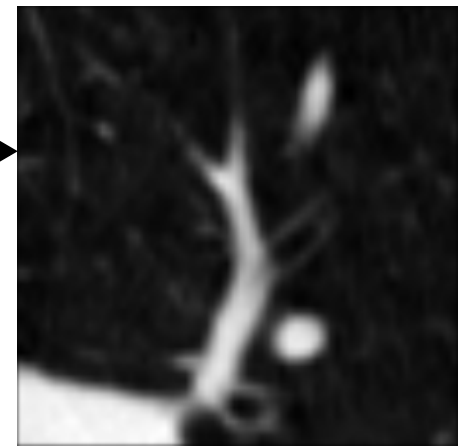
IV. In-painting



Deformed Image



Original Image

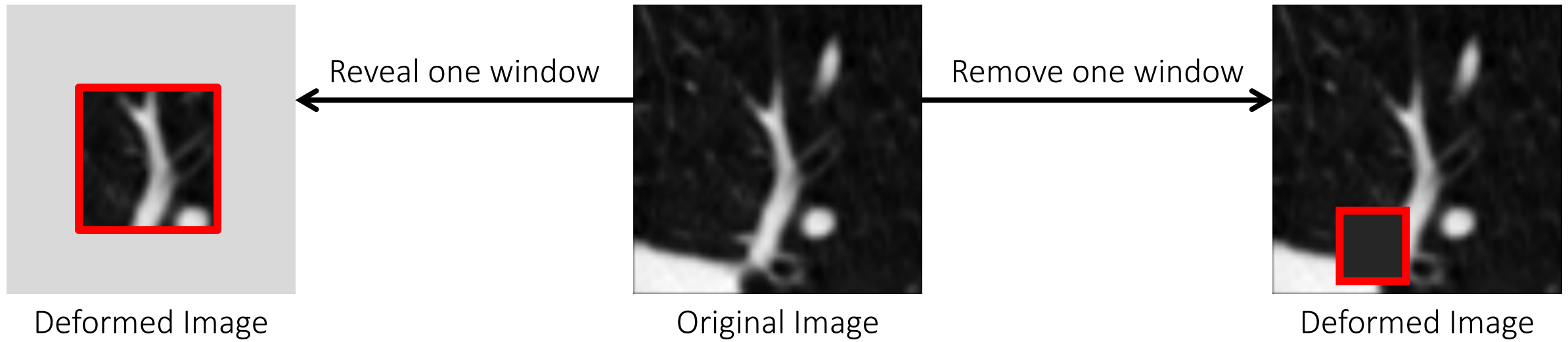


Deformed Image



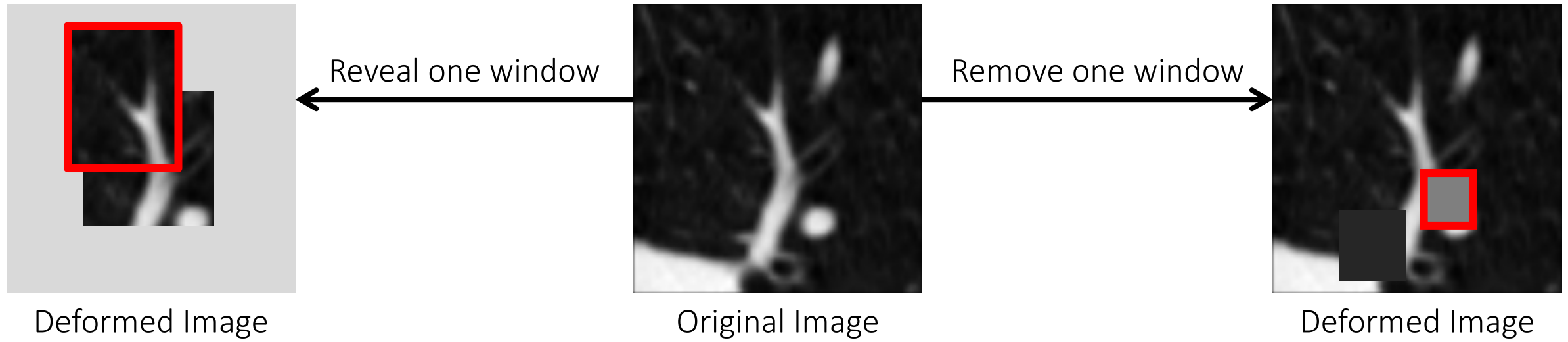
III. Out-painting

IV. In-painting



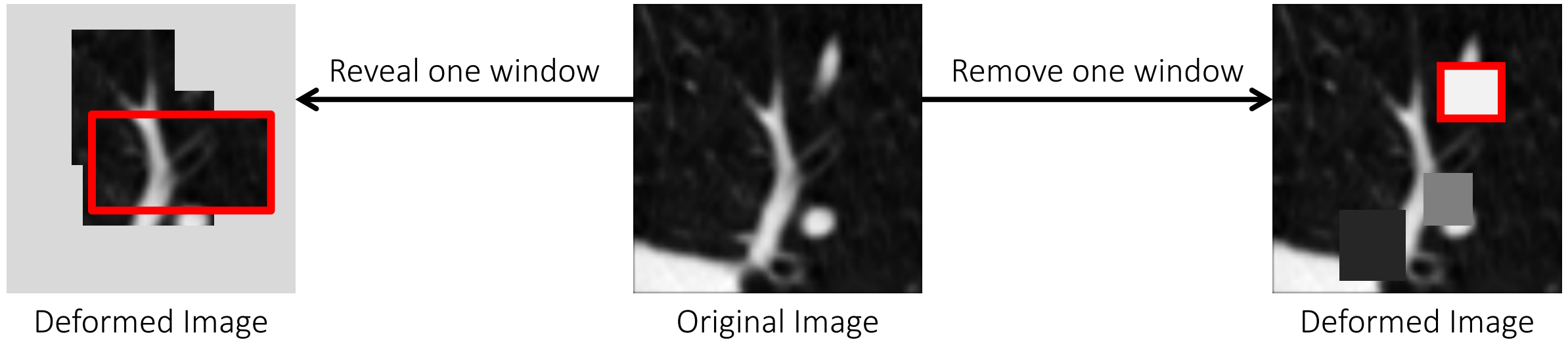
III. Out-painting

IV. In-painting



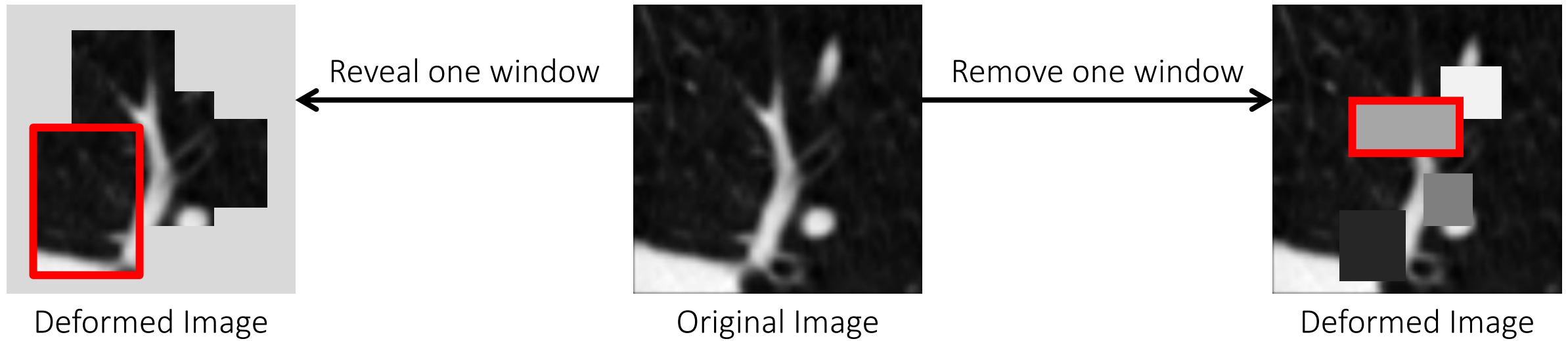
III. Out-painting

IV. In-painting



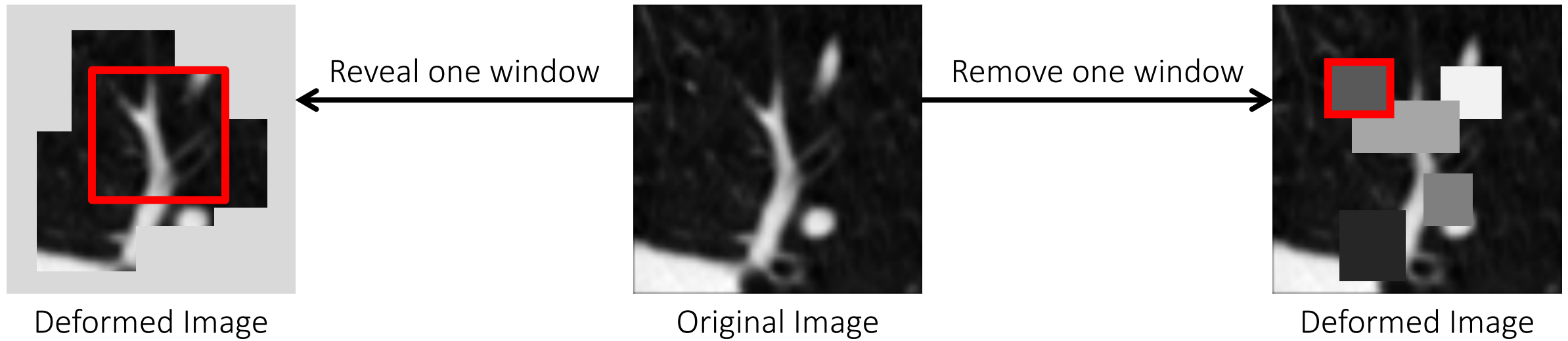
III. Out-painting

IV. In-painting



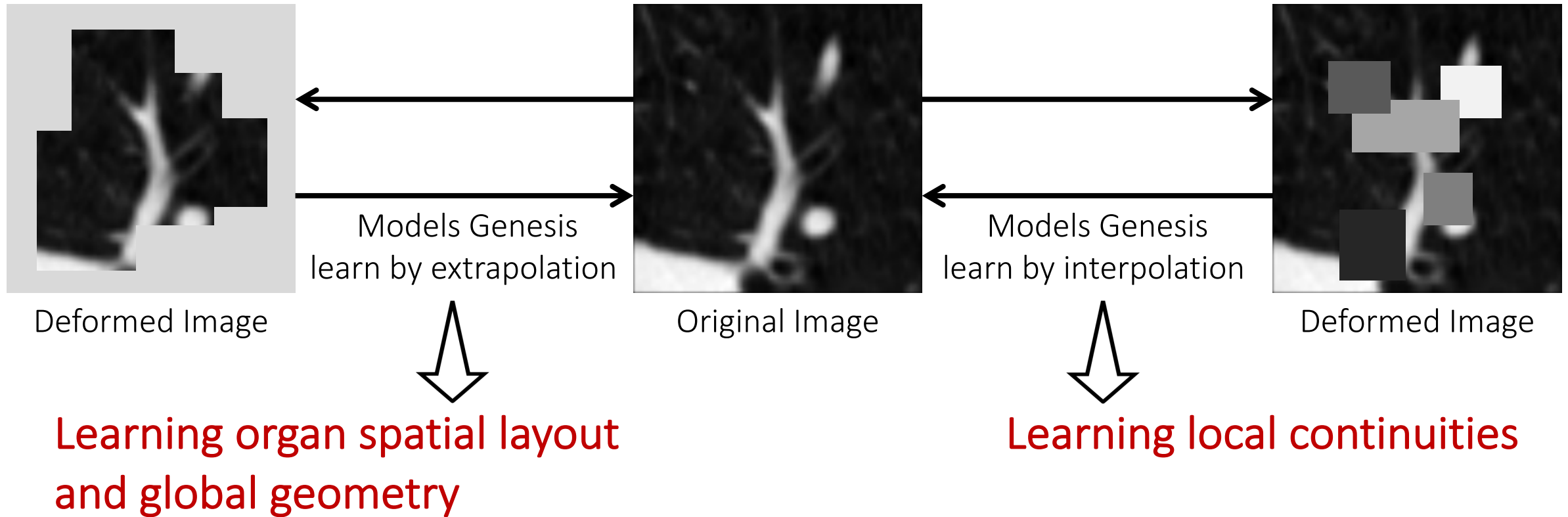
III. Out-painting

IV. In-painting



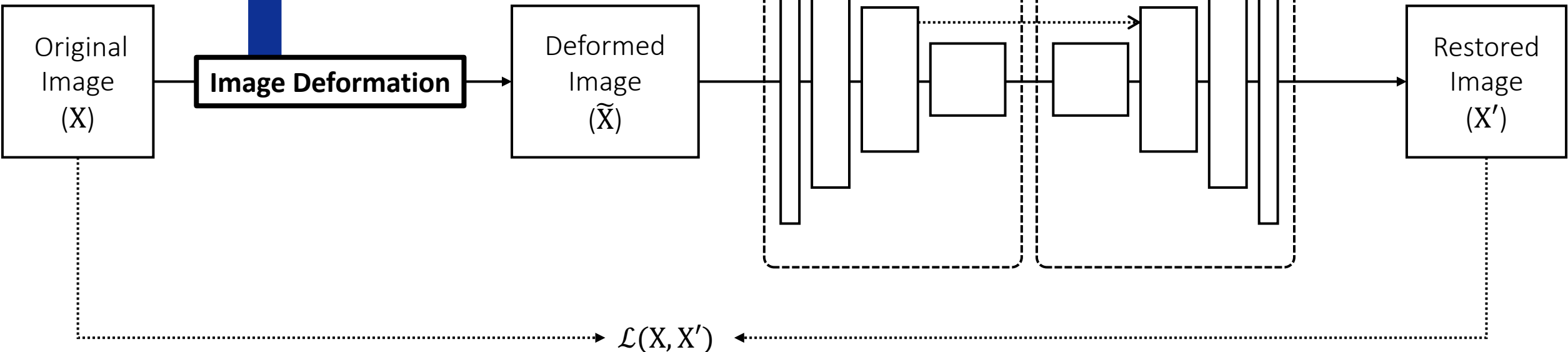
III. Out-painting

IV. In-painting

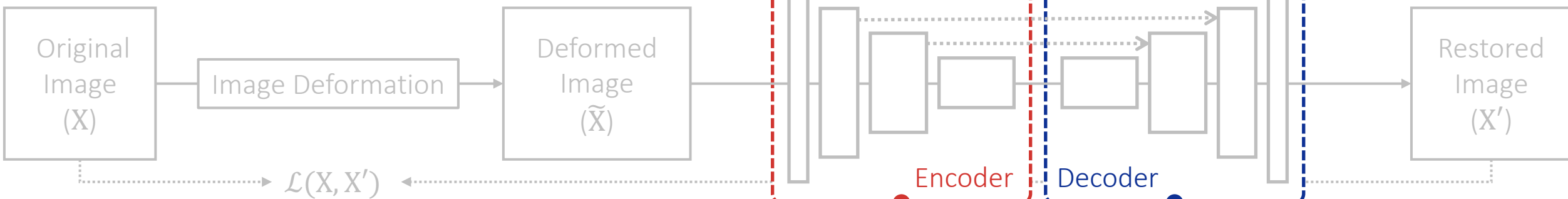
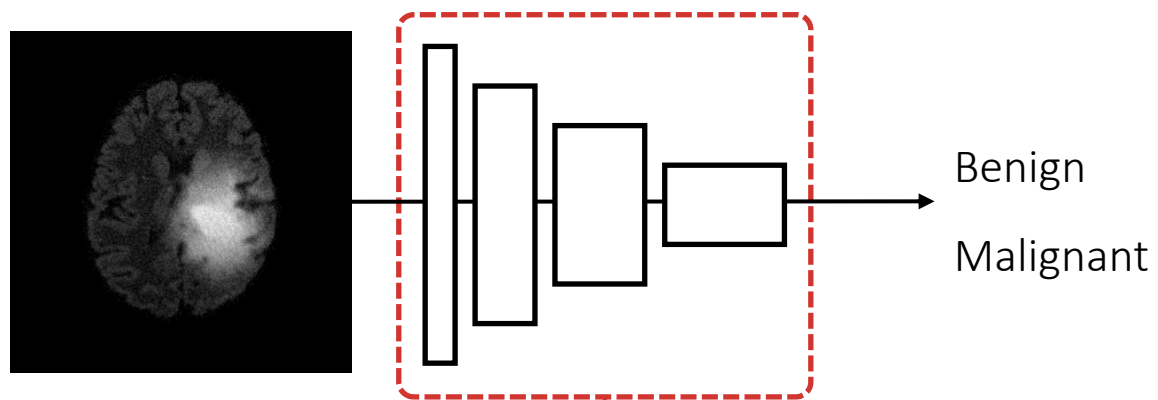


- → Non-linear
- → Local shuffling
- → Out-painting
- → In-painting
- → More ...

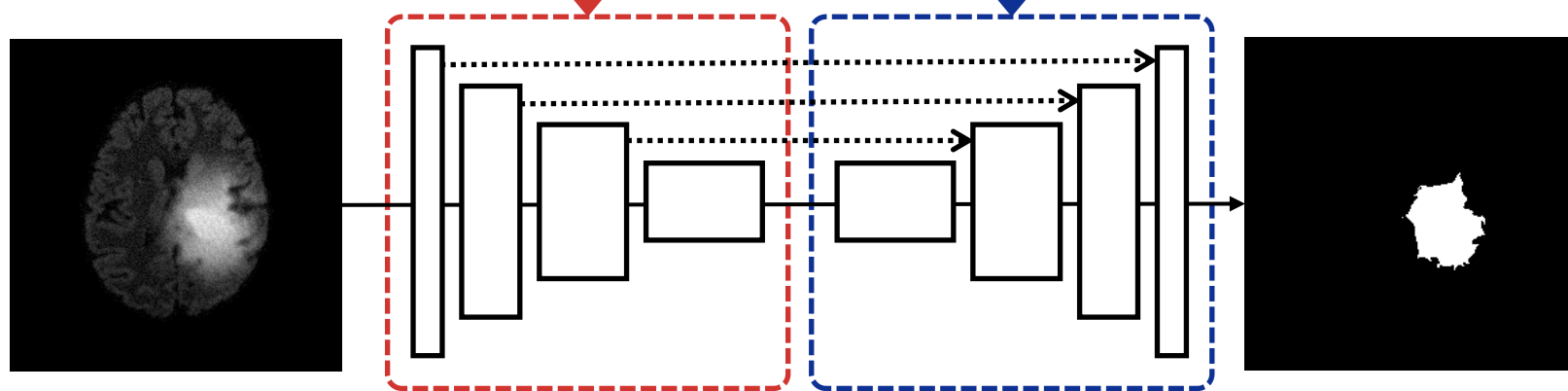
Combination: learning from multiple perspectives
e.g., organ appearance, texture, boundary, global geometry, and local continuity



Once pre-trained,
the **encoder** could be used
for target classification tasks
e.g., brain tumor classification;

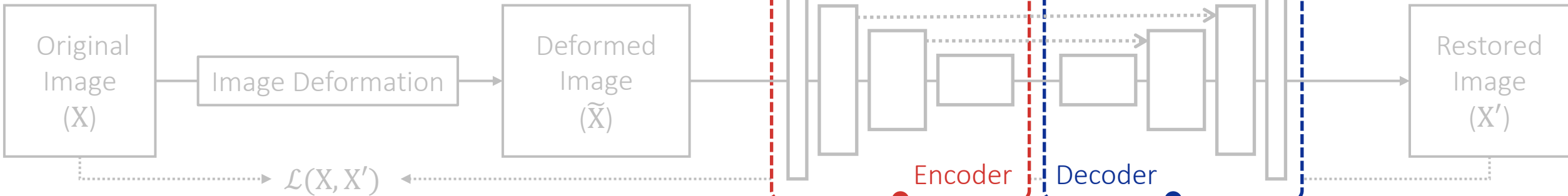


the **encoder-decoder** could be used
for target segmentation tasks
e.g., brain tumor segmentation



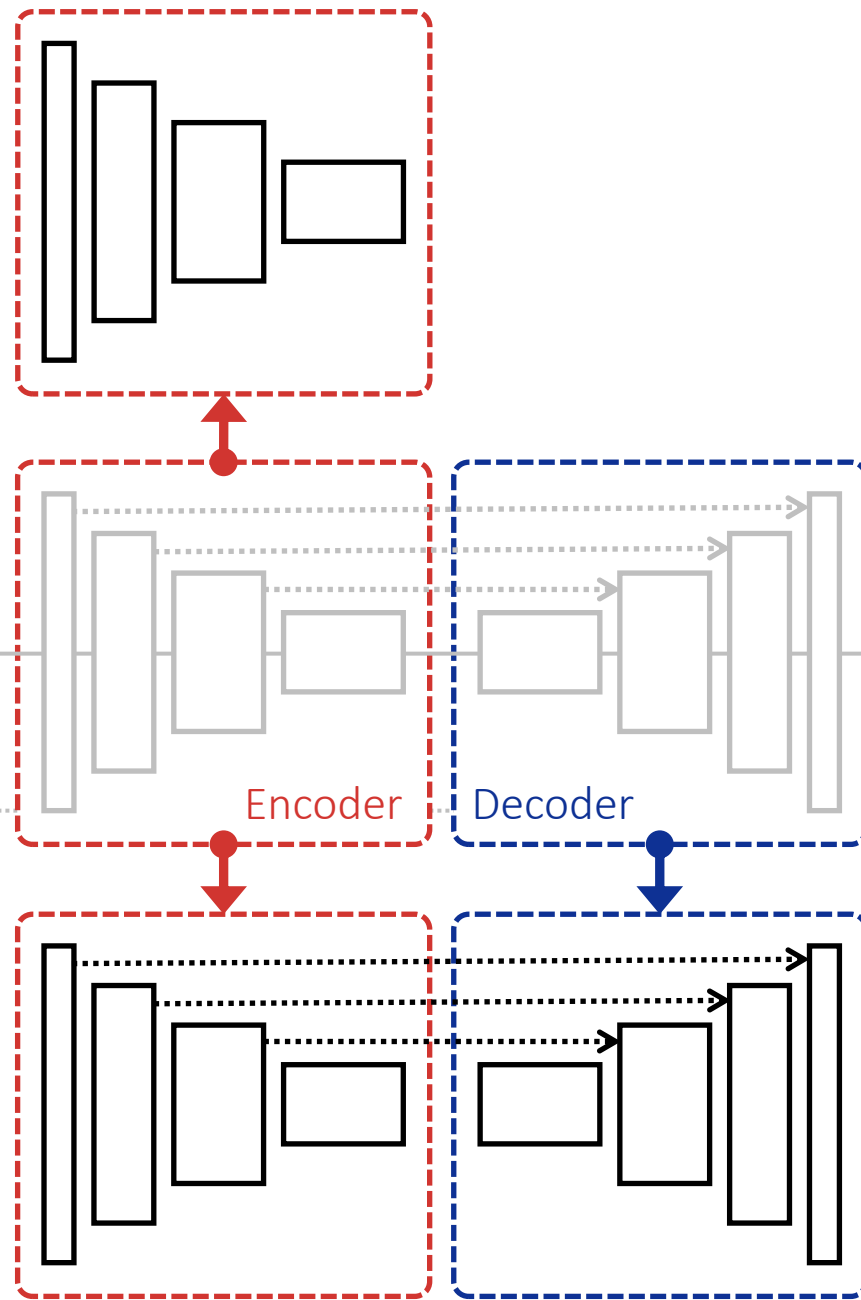
Four classification applications

1. Lung nodule false positive reduction (CT)
2. PE false positive reduction 3D CT (CT)
3. Eight pulmonary diseases classification (X-ray)
4. Rol/bulb/background classification (Ultrasound)



Three segmentation applications

1. Lung nodule segmentation (CT)
2. Liver segmentation (CT)
3. Brain tumor segmentation (MRI)



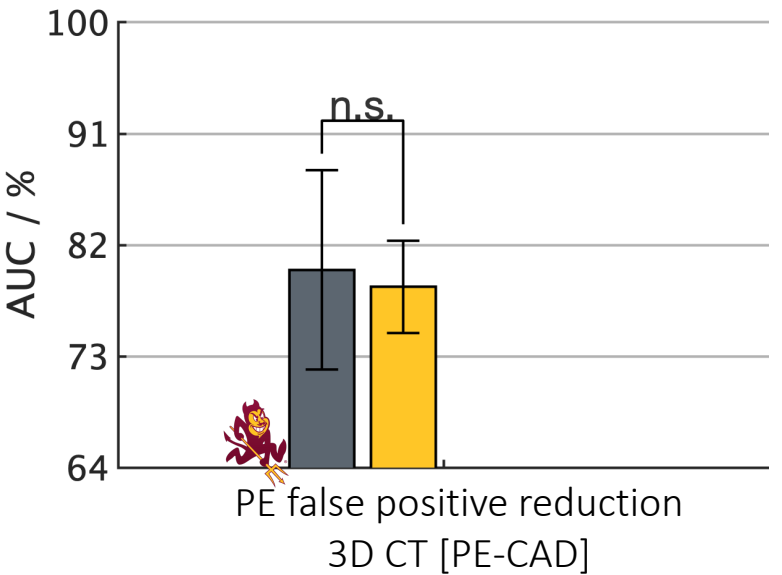
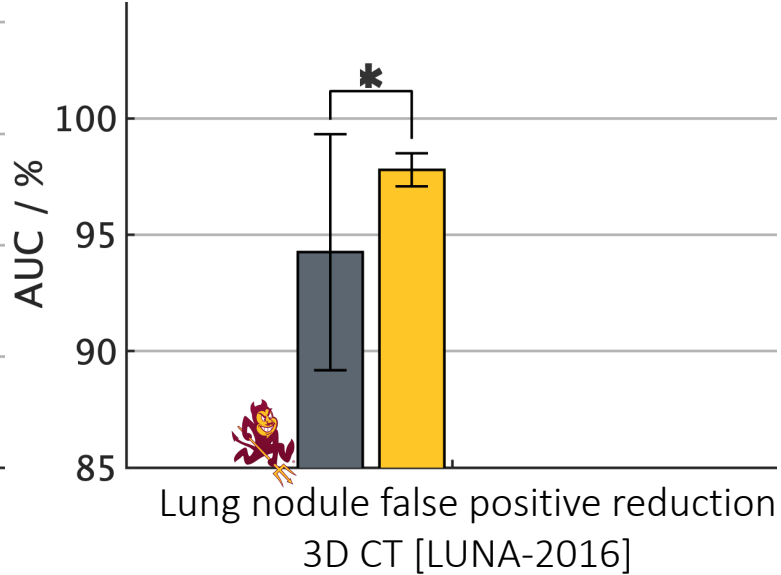
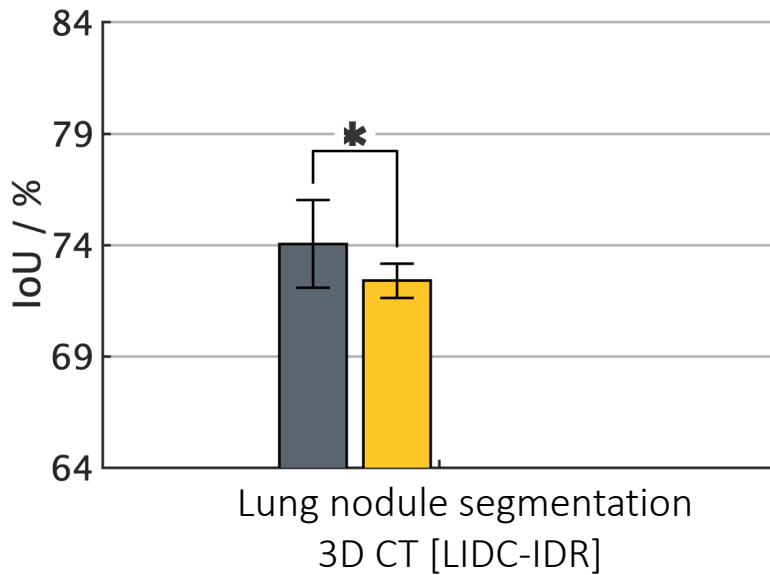
Devils in 3D Models



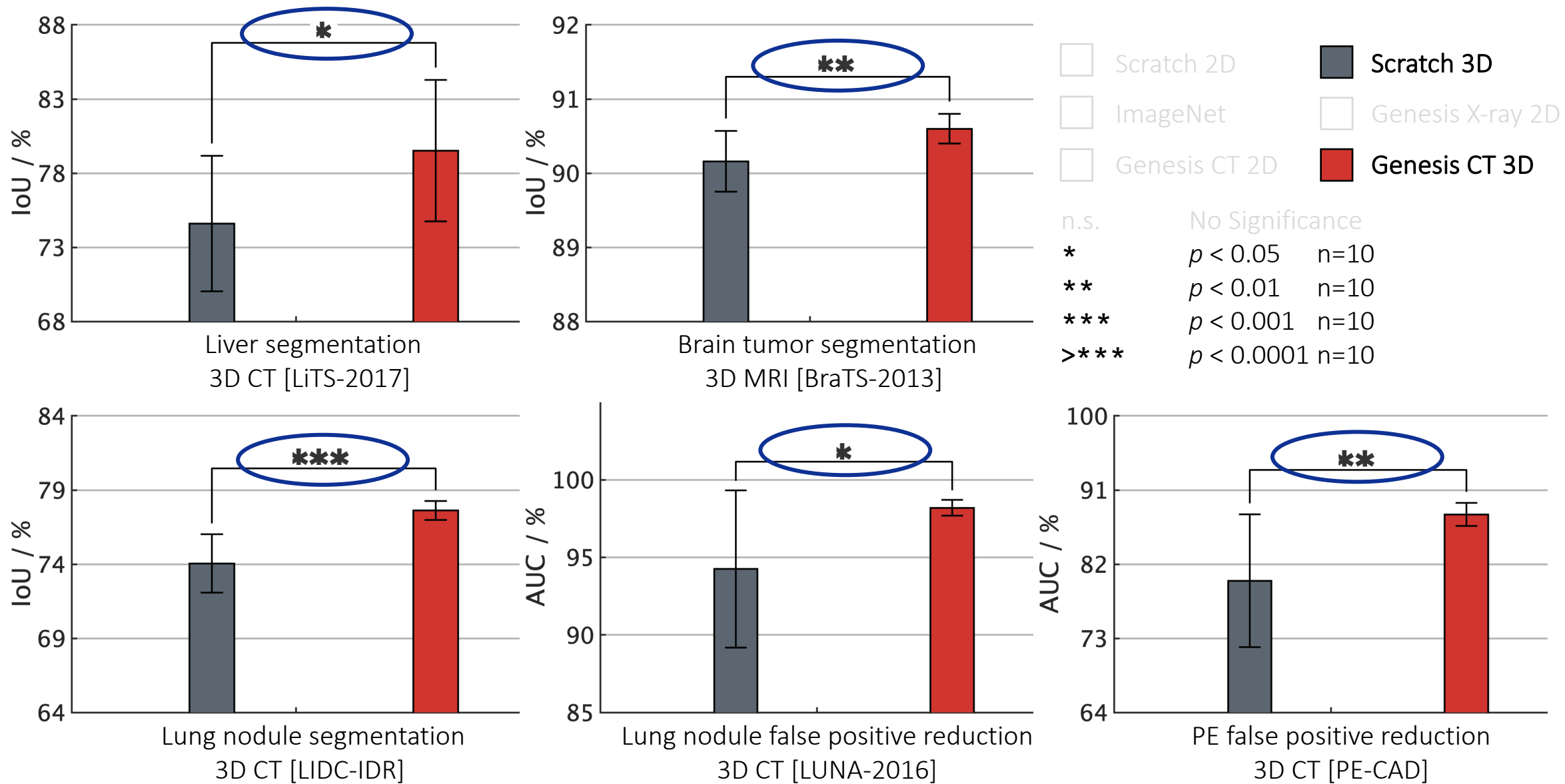
: Learning from scratch *simply* in 3D may not necessarily yield performance better than fine-tuning from ImageNet in 2D



n.s. No Significance
 * $p < 0.05$ n=10
 ** $p < 0.01$ n=10
 *** $p < 0.001$ n=10
 >*** $p < 0.0001$ n=10



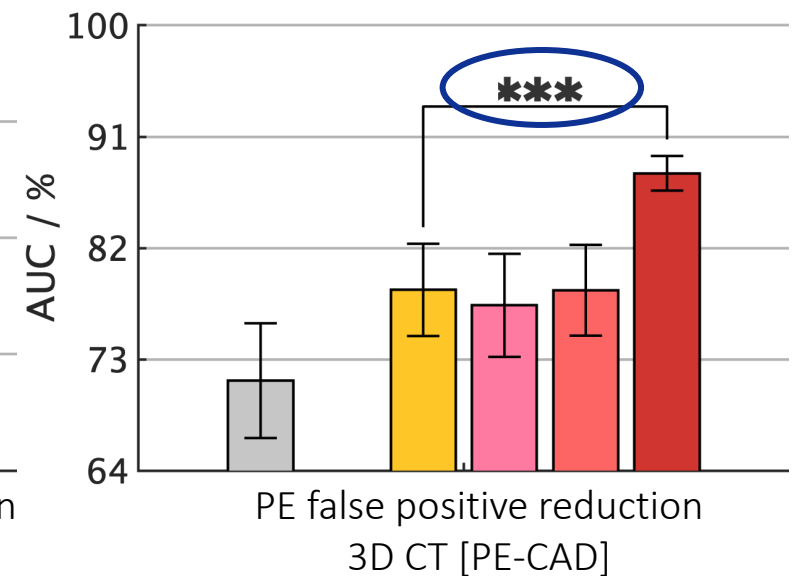
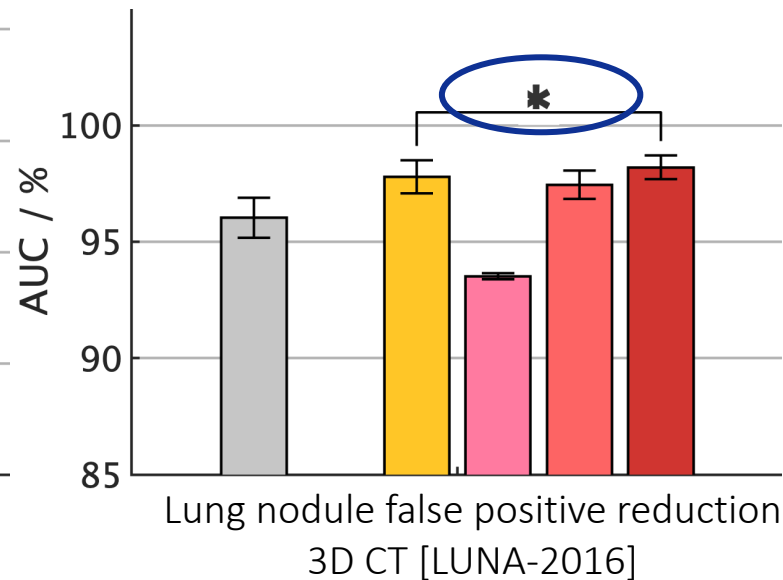
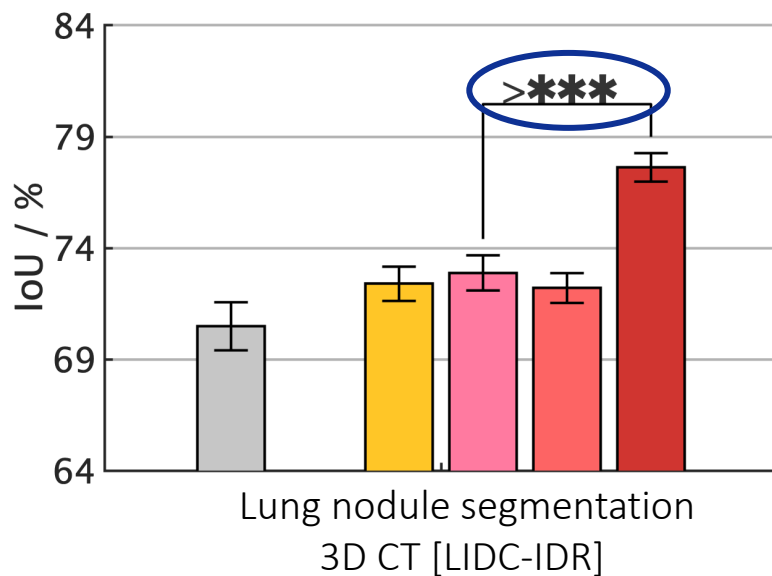
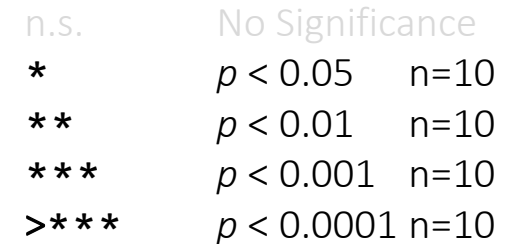
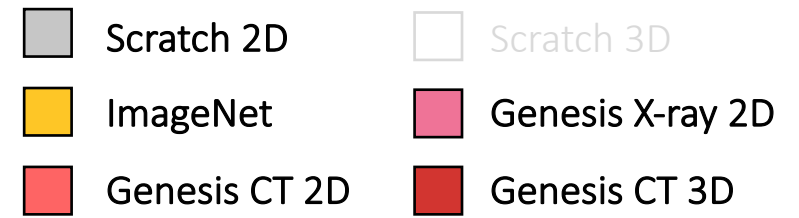
Result I: Models Genesis outperform 3D models trained from scratch



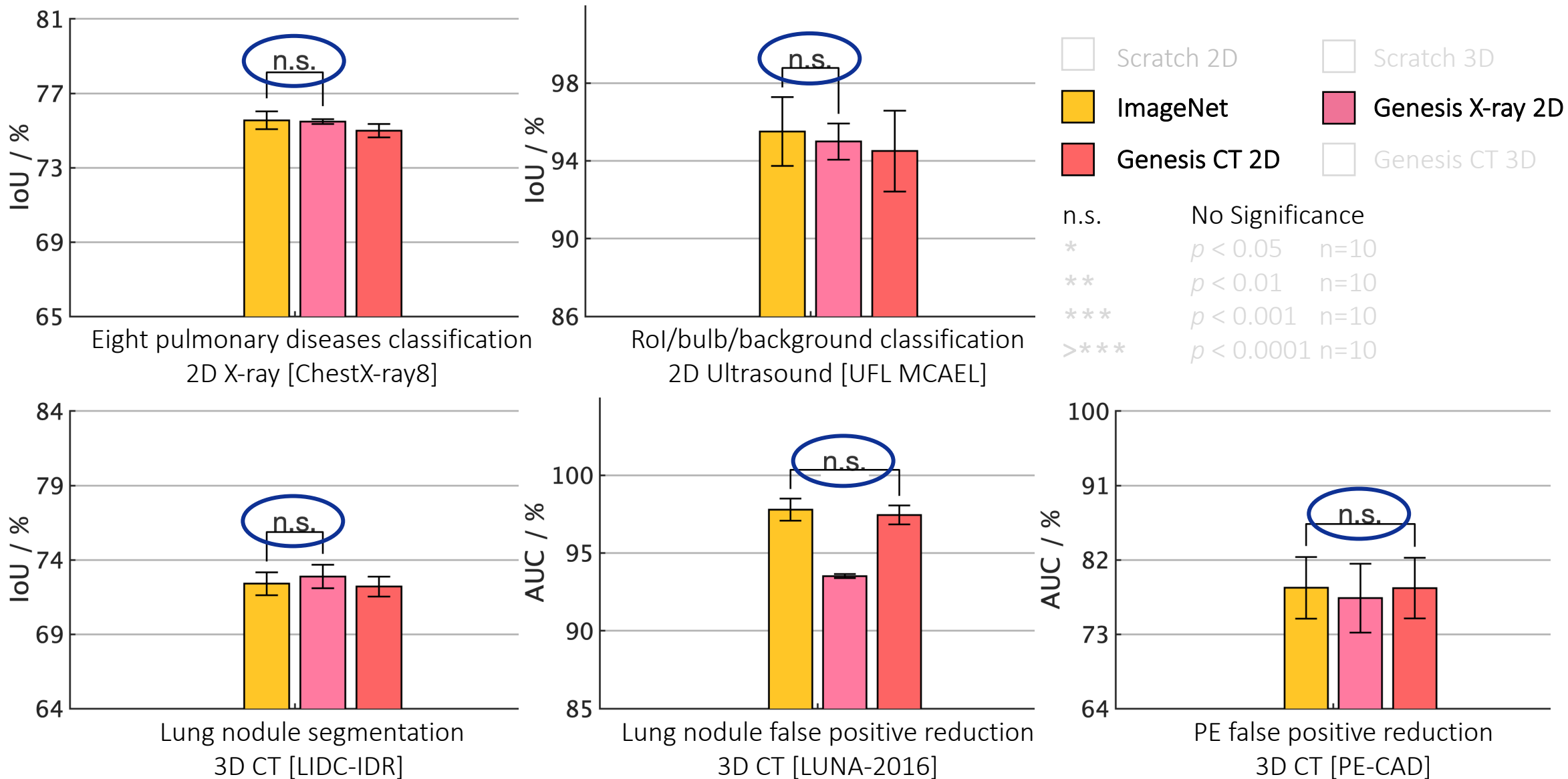
Result II: Models Genesis consistently outperform any 2D approaches

including

1. ImageNet (state-of-the-art)
2. Models Genesis 2D (degraded)
 - Genesis X-ray 2D: pre-trained on NIH X-ray dataset
 - Genesis CT 2D: pre-trained on LUNA-2016 dataset



Result III: Models Genesis 2D (self-supervised) \approx ImageNet (supervised)



Models Genesis: Generic Autodidactic Models for 3D Medical Image Analysis

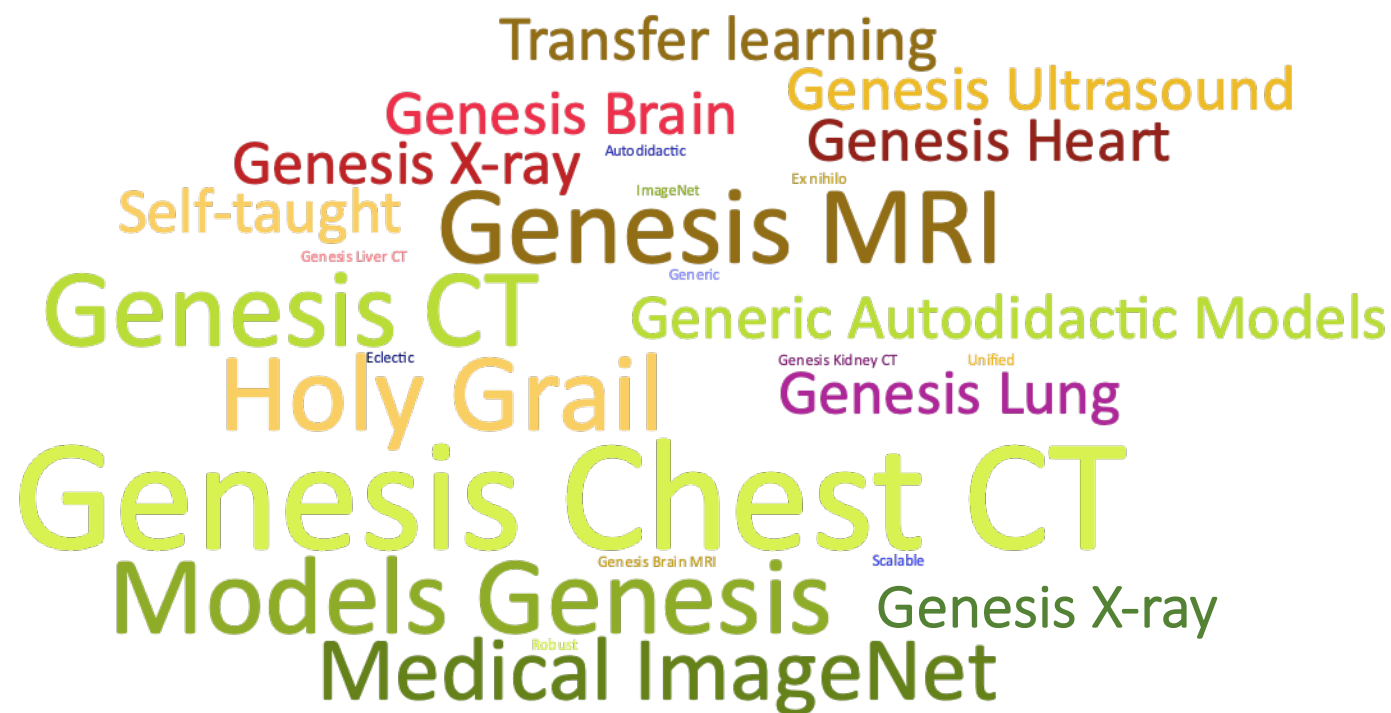
We offer a set of powerful pre-trained 3D models, concluding that

1. Models Genesis outperform 3D models trained from scratch
2. Models Genesis consistently outperform any 2D approaches
3. Models Genesis (2D) offer performances equivalent to supervised pre-trained models

Genesis Chest CT

Genesis X-ray

Models Genesis: Generic Autodidactic Models for 3D Medical Image Analysis



Models Genesis: Generic Autodidactic Models for 3D Medical Image Analysis

Poster session

Tuesday, October 15
13:00 – 14:00

Poster T-5-B-013

Try it for yourself

Code, data, and models
are available online



github.com/MrGiovanni/ModelsGenesis