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# **Domain Adaptation for Medical Image Segmentation on Lung Tumour Datasets**

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# Acronyms

<b>CNN</b>	Convolutional Neural Network
<b>CT</b>	Computed Tomography
<b>DCMQI</b>	DICOM for Quantitative Imaging
<b>DICOM</b>	Digital Imaging and Communications in Medicine
<b>GAN</b>	Generative Adversarial Network
<b>KL</b>	Kullback–Leibler
<b>MRI</b>	Magnetic Resonance Imaging
<b>MSA</b>	Multihead Self-attention
<b>MSD</b>	Medical Segmentation Decathlon
<b>NIfTI</b>	Neuroimaging Informatics Technology Initiative
<b>NLP</b>	Natural Language Processing
<b>NSCLC</b>	Non-Small Cell Lung Cancer
<b>RIDER</b>	The Reference Image Database to Evaluate Therapy Response
<b>SCLC</b>	Small Cell Lung Cancer
<b>TCIA</b>	The Cancer Imaging Archive
<b>UDA</b>	Unsupervised Domain Adaptation
<b>U-Net</b>	Convolutional Networks for Biomedical Image Segmentation
<b>UNETR</b>	UNet Transformer
<b>VAE</b>	Variational Autoencoder
<b>ViT</b>	Vision Transformer

# Abstract

Over the past few decades, advances have been made in the classification, diagnosis and treatment of lung cancer in many ways. Machine learning-based lung cancer prediction models such as semantic segmentation have been proposed to assist clinicians in the management of incidentally detected or screened-out indeterminate lung nodules. Semantic segmentation transforms raw medical images into clinically relevant, spatially structured information, such as outlining tumour boundaries and is an essential prerequisite for abundant clinical applications. However, the normal segmentation models may not have qualified performance, due to the lack of labelled medical imaging datasets. Domain adaptation can be introduced to finetune the target datasets on one pre-trained model for better accuracy. This project focuses on applying a segmentation and shaping model based domain adaptation framework with 3D U-Net and UNETR as its backbone. Experimental results demonstrate that the Dice gap between a certain method and direct tests is around 2% in most cases, where three datasets are imported including MSD, RIDER, and NSCLC. Domain adaptation significantly improves the performance on the small dataset NSCLC of models pre-trained on the large dataset RIDER. In addition, the 3D U-Net with simple network architecture trained on the relatively abundant datasets RIDER reaches the highest scores among the current results.

**Key Words:** Deep Learning, Convolutional Neural Network, Medical Segmentation, Unsupervised Domain Adaptation, Lung Tumour Dataset



# Acknowledge

I am grateful to all of those who have helped me during this project, especially my supervisor Prof. Erick Purwanto.

I would like to thank my parents, my girlfriend and my roommates, although we get together less and leave more.

I also thank PyCharm and my NVIDIA RTX 2060 for spending many lonely nights with me.

# Chapter 1

## Introduction

### 1.1 Motivation, Aims and Objective

Semantic segmentation is a computer vision task that aims to assign a class to each pixel in the image using that image as input [8]. If multiple objects of the same class are accessible, they can be simply labelled with their class [8]. Semantic segmentation has many applications for medical image analysis, such as segmenting pancreas tumour regions in portal venous phase Computed Tomography (CT) scans [8]. This project focuses on some lung cancer datasets which require the participants to annotate the tumour in the lungs, such as Non-Small Cell Lung Cancer (NSCLC), considering large-ranging foreground size as a challenge. NSCLC is any type of epithelial lung cancer other than Small Cell Lung Cancer (SCLC), where the most common types of NSCLC are squamous cell carcinoma, large cell carcinoma, and adenocarcinoma [9, 10]. However, machine learning techniques for computer-aided medical image analysis are often plagued by domain transfer problems caused by different distributions between source and target data [11]. For example, the normal segmentation models may not have qualified performance, due to the lack of labelled medical imaging datasets. This project aims to build a machine-learning model based on principles of domain adaptation to solve the task of segmenting objects of interest in medical images by training on a general dataset. The model can be easily fine-tuned to fit other datasets with a similar task without extra training.

### 1.2 Literature Review

For medical image segmentation, Convolutional Networks for Biomedical Image Segmentation (U-Net) is a widely-used fully convolutional network under the encoder-decoder backbone, which works with very few training images and yields precise segmentations [12]. 3D U-Net is a derivative network developed for sparsely annotated volumetric images, such as CT volume [5]. Furthermore, Zhang et al. [13] introduce Deep Residual U-Net (ResUNet) to accomplish feature accumulation in recursive residual convolution layers based on U-Net, considering the time dependency of image sequences. An adversarial-based method is a promising approach for domain adap-

tation to training robust deep networks by complex samples across diverse domains [14]. The image-to-image translation for converting images from source to target domain can be realised by a Generative Adversarial Network (GAN) [15]. In addition, Liu et al. [16] proved that Variational Autoencoder (VAE) could learn the shape distribution for a specific organ in unsupervised domain adaptation. Synergistic Image and Feature Adaptation (SIFA) is one of unsupervised domain adaptation framework which presents synergistic fusion of adaptations from both image and feature perspectives, guided by adversarial losses [17].

### 1.3 Industrial Relevance

Semantic segmentation transforms raw medical images into clinically relevant, spatially structured information, such as outlining tumour boundaries and is an essential prerequisite for abundant clinical applications, such as radiotherapy planning and treatment response monitoring, and provides new insights into the early diagnosis of the corresponding disease [18]. For example, early detection of abnormal signs of diabetic retinopathy can lead to effective treatment before its initial onset and prevent blindness in more than 50% of cases [19]. However, manual segmentation of medical images for tissues such as retinal blood vessels is a lengthy and tedious task, requiring extra training and professional skill [20]. Furthermore, deep learning networks adaptable for a particular clinical problem may not necessarily generalise well to different, unexplored tasks [18]. Unsupervised domain adaptation can be seen as an approach for image segmentation by some labelled data without human intervention. This method would enhance the technical scalability to allow many new applications in computer-aided diagnosis, biomarker extraction, surgical intervention planning, disease prognosis, etc. [18].

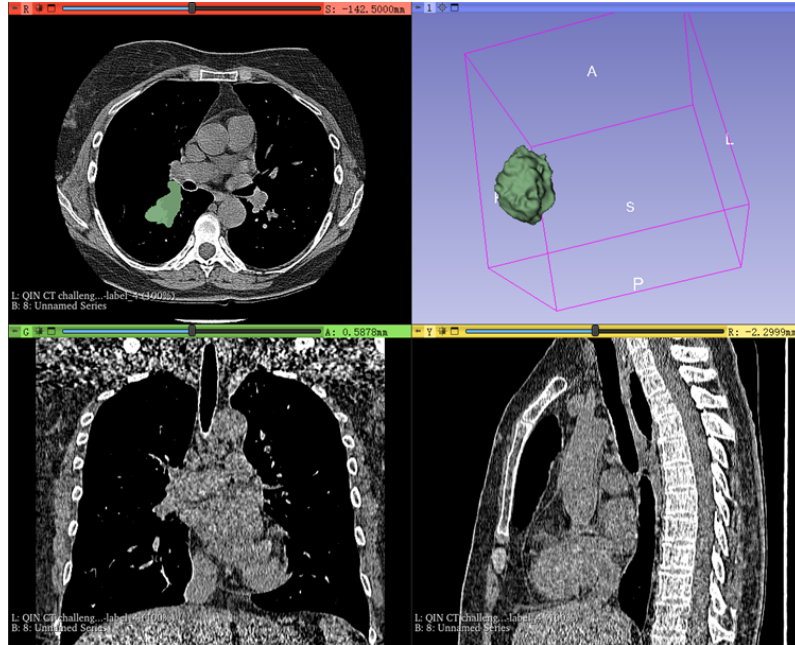


Fig. 1.1: The model of one CT set in RIDER Lung CT displayed in 3D Slicer.

# Chapter 2

## Datasets

### 2.1 Dataset Introduction

For domain adaptation, the variation of preferred modality and scanning protocol should be considered to improve the network’s versatility. The Cancer Imaging Archive (TCIA) [21] is identified as the primary dataset source for retrieving data for this project to avoid data conflicts. This project selects a subset of the Non-Small Cell Lung Cancer (NSCLC) dataset [1], The Reference Image Database to Evaluate Therapy Response (RIDER) lung CT dataset [22], as the source dataset to train the source network, considering its data size and data diversity, Another sub-dataset of NSCLC, NSCLC Radiomics Interobserver1 [23], and Lung Tumours dataset from Medical Segmentation Decathlon (MSD) [18] are selected as the transfer data to finetune the target network.

#### 2.1.1 Medical Segmentation Decathlon

This Medical Segmentation Decathlon (MSD) challenge and dataset aims to provide such resource through the open sourcing of large medical imaging datasets on several highly different tasks, and by standardising the analysis and validation process [18].

#### Lung Tumours

The dataset consists of preoperative thin-section CT scans from 96 patients with non-small cell lung cancer. This data set was selected due to the challenge of segmenting small tumour regions in an image with a large field-of-view [18]. The dataset consists of training set, testing set, and label set.

1. For training set, it includes 64 CT NIfTI files.
2. For testing set, it includes 32 CT NIfTI files.
3. For label set, it includes 64 segmentation NIfTI files.

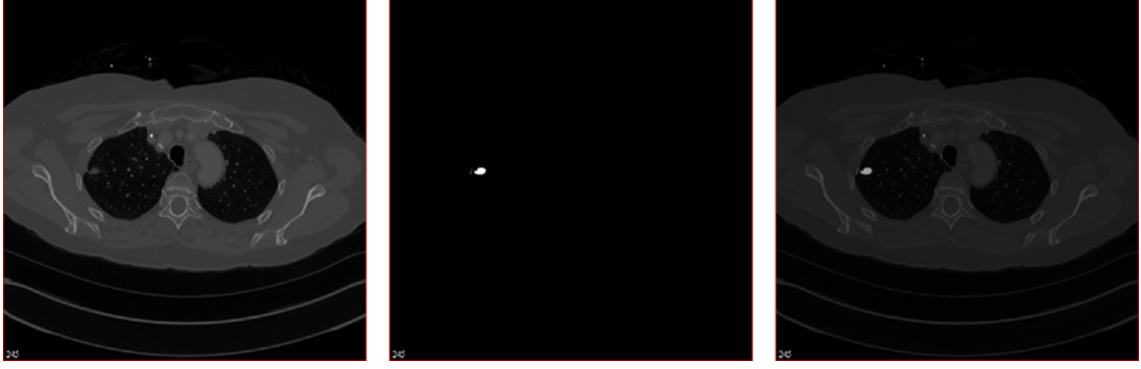


Fig. 2.1: The training, label, and composite CT image of "lung\_001.nii.gz" in MSD lung tumour dataset at frame 245.

### 2.1.2 Non-Small Cell Lung Cancer

The Non-Small Cell Lung Cancer (NSCLC) dataset is published in Nature Communications which applies a radiomic approach to computed tomography data of 1,019 patients with lung or head-and-neck cancer [1]. Radiomics refers to the comprehensive quantification of tumour phenotypes by applying a large number of quantitative image features [1]. This project introduces two lung tumour relevant datasets through manual delineation of the 3D volume of the tumor.

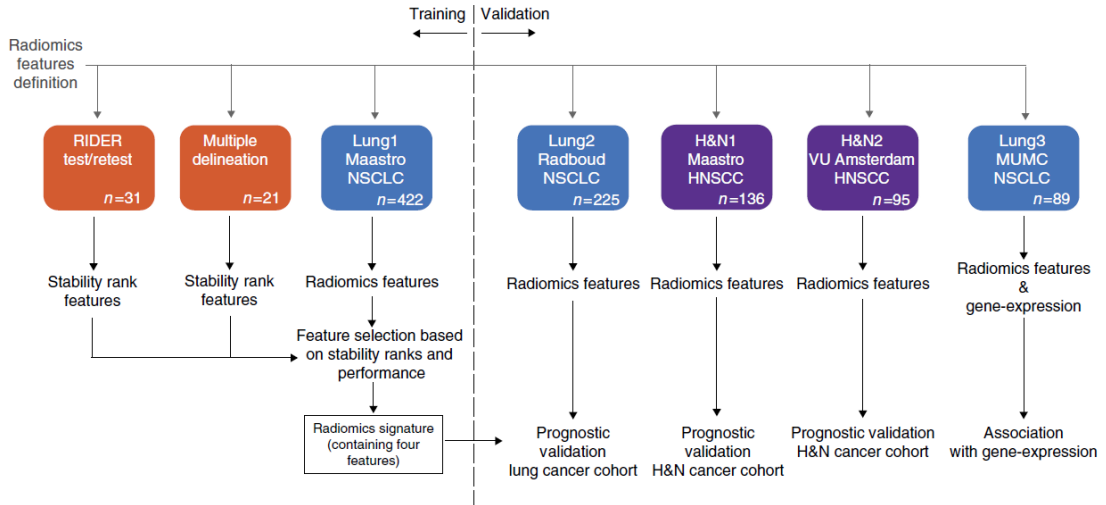


Fig. 2.2: Analysis workflow of this datasets collection which includes RIDER datasets and NSCLC Radiomics datasets [1].

### RIDER Lung CT

The Reference Image Database to Evaluate Therapy Response (RIDER) Lung CT collection was constructed as part of a study to evaluate the variability of tumor unidimensional, bidimensional, and volumetric measurements on same-day repeat CT scans in patients with non-small cell lung cancer [22]. The dataset consists of raw resources (RIDER Lung CT) [22] and third party analyses

(RIDER-LungCT-Seg) [24].

1. For RIDER Lung CT, it includes 63 CT DICOM series from 63 patients.
2. For RIDER-LungCT-Seg, it includes 59 relevant SEG and RESTRUCT DICOM series.

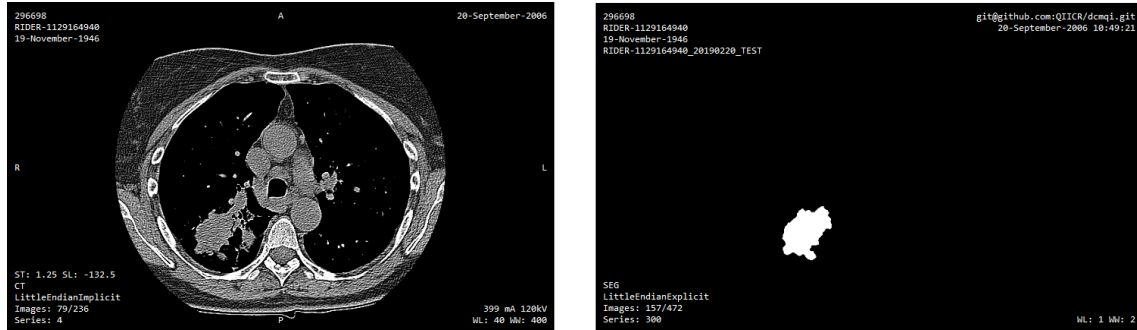


Fig. 2.3: The raw and segmentation CT image of "RIDER-1129164940" in RIDER lung tumour dataset at frame 157.

### NSCLC-Radiomics-Interobserver1

This collection contains clinical data and CT from 22 non-small cell lung cancer radiotherapy patients [23]. The dataset consists of CT and segmentation as well.

1. For CT, it includes 22 CT DICOM series from 22 patients.
2. For segmentation, it includes 22 relevant SEG and RESTRUCT DICOM series.

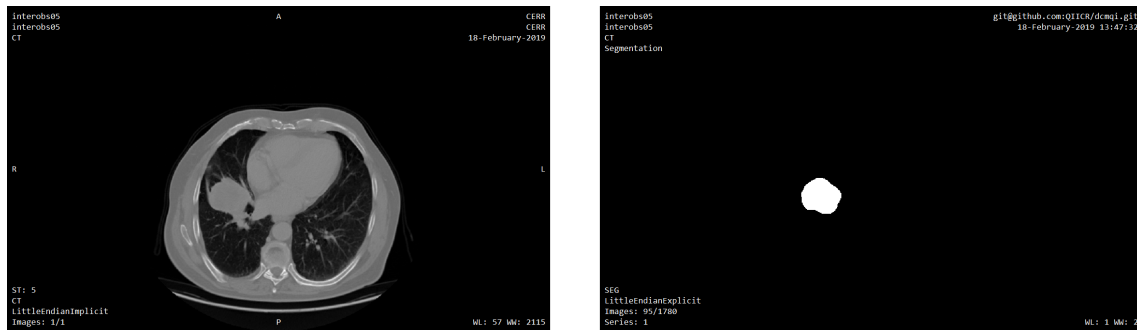


Fig. 2.4: The raw and segmentation CT image of "interobs05" in NSCLC Radiomics Interobserver1 lung tumour dataset at frame 95.

## 2.2 Dataset Format

### 2.2.1 Neuroimaging Informatics Technology Initiative

Neuroimaging Informatics Technology Initiative (NIfTI) is a new Analyze-style data format as a short-term measure to facilitate inter-operation of functional MRI data analysis software packages [25]. The original format of MSD datasets is NIfTI.

```

>>> import nibabel as nib
>>> path = r"dataset/nih_data/Task06_Lung/imagesTr/lung_001.nii.gz"
>>> image = nib.load(path)
>>> type(image.get_fdata())
<class 'numpy.ndarray'>
>>> image.get_fdata().shape
(512, 512, 304)

```

Fig. 2.5: The NIfTI file loading process of "lung\_001.nii.gz" in MSD lung tumour dataset.

## 2.2.2 Digital Imaging and Communications in Medicine

Digital Imaging and Communications in Medicine (DICOM) is the international standard for medical images and related information [26]. The raw CT radiomic images in the NSCLC datasets are all in the DICOM format. DICOM for Quantitative Imaging (DCMQI) is a protocol with minimum dependencies to support standardized communication of quantitative image analysis research data using DICOM standard [27]. The segmentation slices in the NSCLC datasets are all in the DCMQI format.

## 2.2.3 Medical Format Normalization

Considering the less metadata and better performance in 3D volumn representation, this project prefers to use NIfTI as an intermediate format towards NumPy array. NumPy arrays as a standard matrix format provide more transformation functions and better compatibility with PyTorch. The specific reformatting path is depicted in the Fig 2.6, where some third-party tools are introduced, such as dcm2niix and dcmqi.

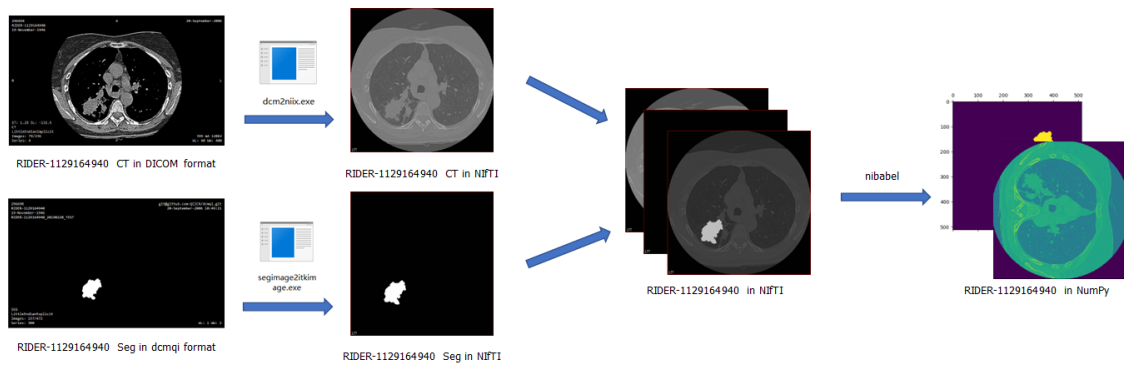


Fig. 2.6: The files reformatting process from DICOM and DCMQI to NIfTI and finally NumPy arrays.

## 2.3 Dataset Preprocess

### 2.3.1 Resampling

Due to the different voxel coordinates of CT and segmentation series, the CT and segmentation volumes are resampled to the unit voxel size of  $1mm \times 1mm \times 1mm$ . The resampling transformation is applied by matrix multiplication with the affine matrix, which is defined as

$$\begin{bmatrix} x \\ y \\ z \\ 1 \end{bmatrix} = \begin{bmatrix} m_{1,1} & m_{1,2} & m_{1,3} & a \\ m_{2,1} & m_{2,2} & m_{2,3} & b \\ m_{3,1} & m_{3,2} & m_{3,3} & c \\ 0 & 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} i \\ j \\ k \\ 1 \end{bmatrix} \quad (2.1)$$

### 2.3.2 Cube Bounding

Cube bounding is utilized to crop the CT and segmentation volumes to emphasize the tumour area and reduce unwanted background. The cube box is centered on the center of the tumour volumes, with enough side length and fixed pads to cover the whole target volumes and necessary backgrounds.

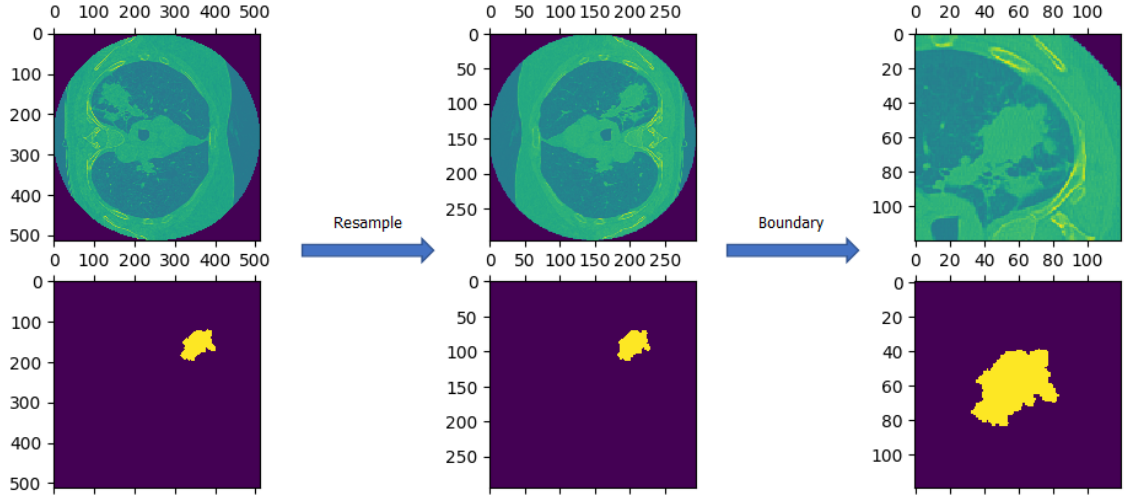


Fig. 2.7: The data preprocessing process through resampling and cube bounding.

## 2.4 Data Augmentation

Data augmentation is often used to handle the problem of the data shortage and insufficient training samples, especially for 3D segmented medical images. In model fitting, it can alleviate the overfitting problem for models with strong generalization capabilities and robustness [28]. This project applies geometric methods for this process, including scaling, rotation and cropping in a



random range. The volumes are through normalization mapped to  $[0, 1]$  at beginning and centered in the patch volumes in the end to standardize the volume size and range.

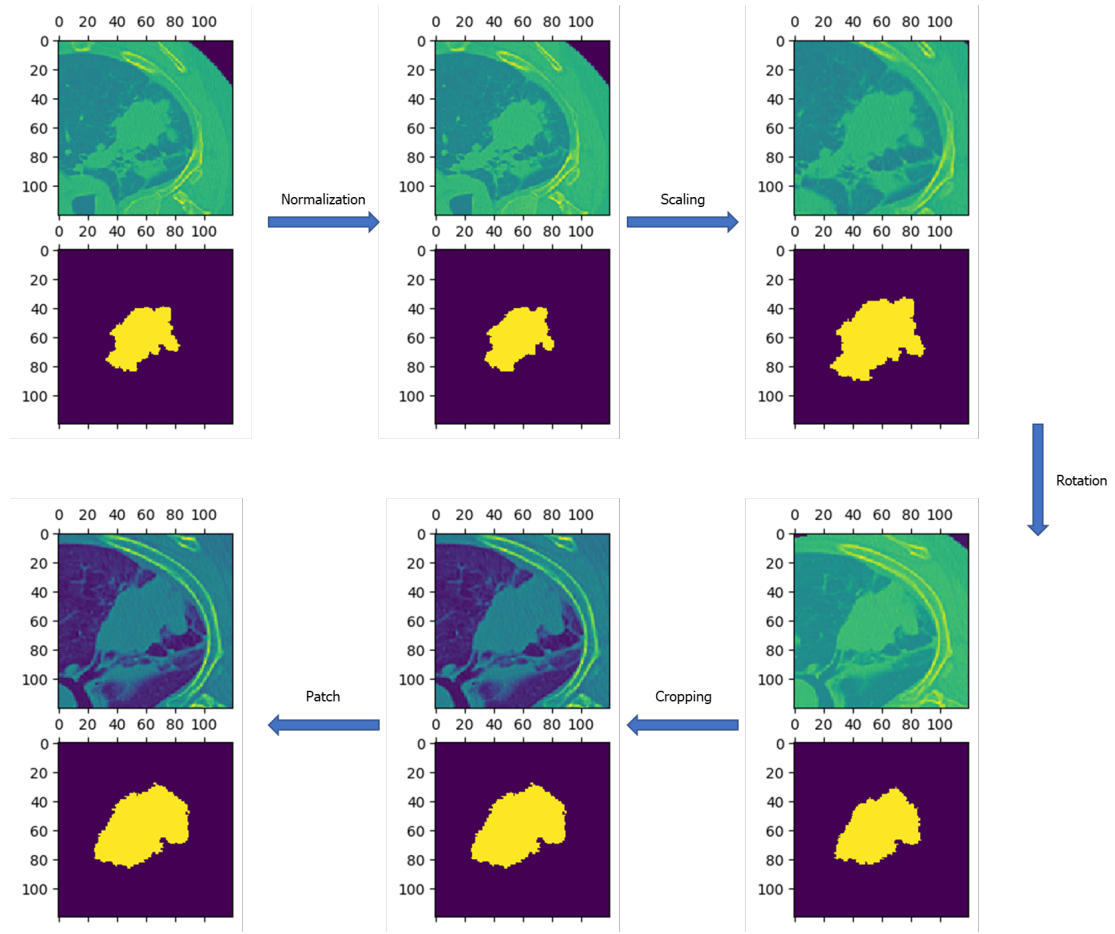


Fig. 2.8: The data augmentation process through normalization, scaling, rotation, cropping and patch.

# Chapter 3

## Methodology

### 3.1 Method Background

#### 3.1.1 3D Convolutional Neural Network

Convolutional Neural Network (CNN) is a neural network primarily applied on images. 3D CNN is constructed by convolutional layers convolving 3D kernels to the cube formed by stacking multiple-dimensional spatial features, compared with 2D CNN, where convolutions are applied on the 2D feature maps to compute features from the single layers only [2]. According to Ji et al. [2], the formula of the value at position  $(x, y, z)$  on the  $j$ th feature map in the  $i$ th layer is given by

$$v_{ij}^{xyz} = \tanh \left( b_{ij} + \sum_m \sum_{p=0}^{P_i-1} \sum_{q=0}^{Q_i-1} \sum_{r=0}^{R_i-1} w_{ijm}^{pqr} v_{(i-1)m}^{(x+p)(y+q)(z+r)} \right), \quad (3.1)$$

where  $(P_i, Q_i, R_i)$  is the size of the 3D kernel, and  $w_{ijm}^{pqr}$  is the  $(p, q, r)$ th value of the kernel connected to the  $m$ th feature map in the previous layer.

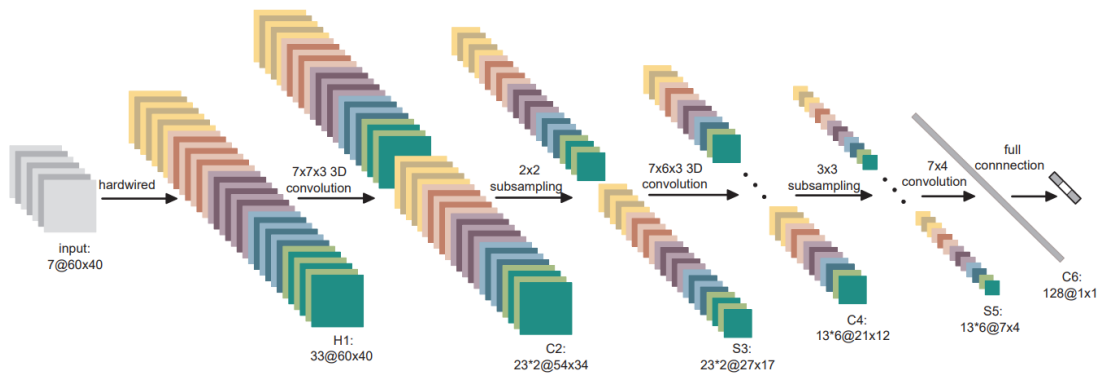


Fig. 3.1: An example 3D CNN architecture for human action recognition Ji et al. [2].

In this project, 3D CNN serves as the basic network block for feature extraction. With relevant de-convolutional networks, segmentation mask can be constructed in image segmentation task.

### 3.1.2 Vision Transformer

Self-attention-based architectures, such as Transformers, have been widely used in the field of Natural Language Processing (NLP). In image recognition, Dosovitskiy et al. [3] introduce Vision Transformer (ViT) architecture which abandons the traditional CNN structure and applies a standard Transformer directly to images. The images are divided into patches linked with the linear sequence, which are seen as tokens in term of NLP [3].

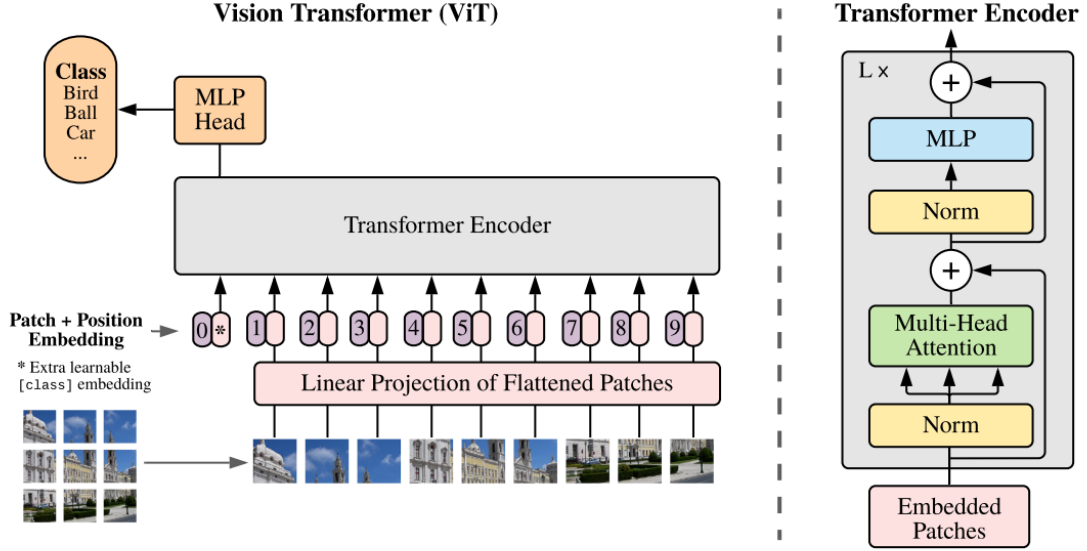


Fig. 3.2: The Vision Transformer architecture [3].

One transformer encoder process can be defined as

$$\begin{aligned}
 \mathbf{z}_0 &= [\mathbf{x}_{\text{class}}; \mathbf{x}_p^1 \mathbf{E}; \mathbf{x}_p^2 \mathbf{E}; \dots; \mathbf{x}_p^N \mathbf{E}] + \mathbf{E}_{\text{pos}}, \quad \mathbf{E} \in \mathbb{R}^{(P^2 \cdot C) \times D}, \mathbf{E}_{\text{pos}} \in \mathbb{R}^{(N+1) \times D} \\
 \mathbf{z}'_\ell &= \text{MSA}(\text{LN}(\mathbf{z}_{\ell-1})) + \mathbf{z}_{\ell-1}, \quad \ell = 1 \dots L \\
 \mathbf{z}_\ell &= \text{MLP}(\text{LN}(\mathbf{z}'_\ell)) + \mathbf{z}'_\ell, \quad \ell = 1 \dots L \\
 \mathbf{y} &= \text{LN}(\mathbf{z}_L^0)
 \end{aligned} \tag{3.2}$$

where MLP is the multilayer perceptron block, MSA is the Multihead Self-attention block, and Layernorm (LN) and residual connection are applied between each network blocks [3].

Multihead Self-attention (MSA) is an extension of standard  $QKV$  self-attention with  $k$  paralleled self-attention operations [3]. The MSA block process can be defined as

$$\begin{aligned}
[\mathbf{q}, \mathbf{k}, \mathbf{v}] &= \mathbf{z} \mathbf{U}_{qkv} & \mathbf{U}_{qkv} &\in \mathbb{R}^{D \times 3D_h} \\
A &= \text{softmax}(\mathbf{q} \mathbf{k}^\top / \sqrt{D_h}) & A &\in \mathbb{R}^{N \times N} \\
SA(\mathbf{z}) &= A \mathbf{v} \\
MSA(\mathbf{z}) &= [SA_1(z); SA_2(z); \dots; SA_k(z)] \mathbf{U}_{msa} & \mathbf{U}_{msa} &\in \mathbb{R}^{k \cdot D_h \times D}
\end{aligned} \tag{3.3}$$

where  $q, k, v$  are query, key and value matrix [3].

Although transformers lack some of the inductive biases inherent to CNNs leading to its poor generalization performance on insufficient amounts of data, some researches, such as UNet Transformer (UNETR) [6] and TransUNet [29], show reasonable segmentation scores by combination of CNNs and transformers [3].

## 3.2 Main Framework

### 3.2.1 VAE Pipeline

Yao et al. [4] developed an Unsupervised Domain Adaptation (UDA) framework to import inherent shape statistics into a standard medical image segmentation model, which is based on a teacher-student learning paradigm with a dual-loss function.

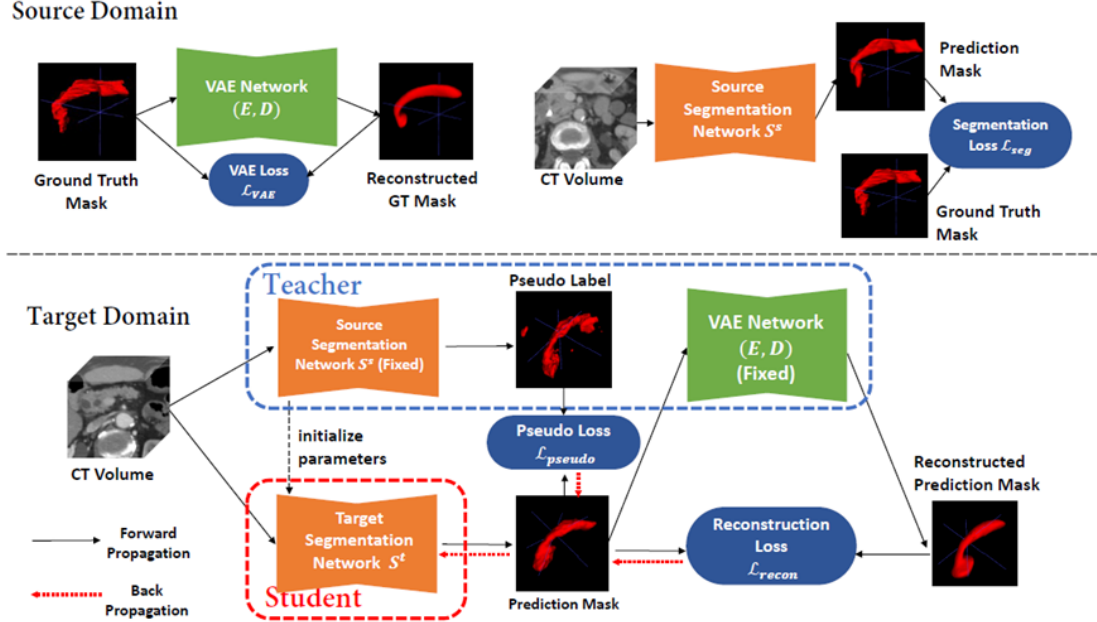


Fig. 3.3: Proposed VAE-based pipeline of Unsupervised Domain Adaptation [4].

On the source domain, the organ shape information is extracted through a pre-trained VAE network in ground truth masks [4]. One dataset is under the training of source segmentation network (3D

U-Net in this project) to initialise it as a teacher network. On the target domain, the two trained networks, i.e., the VAE network and source segmentation network, are fixed as teacher networks [4]. The target segmentation network is initialised by the source one and loads another unlabeled dataset in a different style from the first dataset for domain adaptation [4]. The pseudo label is generated by the source segmentation network in the teacher group for a pseudo-loss calculation to fine-tune the segmentation labels [4]. The distribution of shape for a specific organ is from the VAE network in the teacher group to predict reconstructed masks and acquire reconstruction loss [4]. The dual loss is combined by a loss function with a hyperparameter offset  $\lambda$ , defined in Equation 3.4, considering their adverse effects [4].

$$L_{\theta}(S^t, x^t) = \lambda_{\text{recon}} \cdot \mathcal{L}_{\text{recon}} + \mathcal{L}_{\text{pseudo}} \quad (3.4)$$

### 3.3 Segmentation Network

#### 3.3.1 3D U-Net

3D U-Net is a CNN-based network for image segmentation which learns from sparsely annotated volumetric images [5]. The network extends the previous U-Net architecture from Ronneberger et al. [12] by replacing all 2D operations in dimensions with 3D counterparts [5]. Similar to the standard U-Net, this network contains two paths, including an analysis one and a synthesis one, each with four resolution layers, depicted in Fig 3.4 [5].

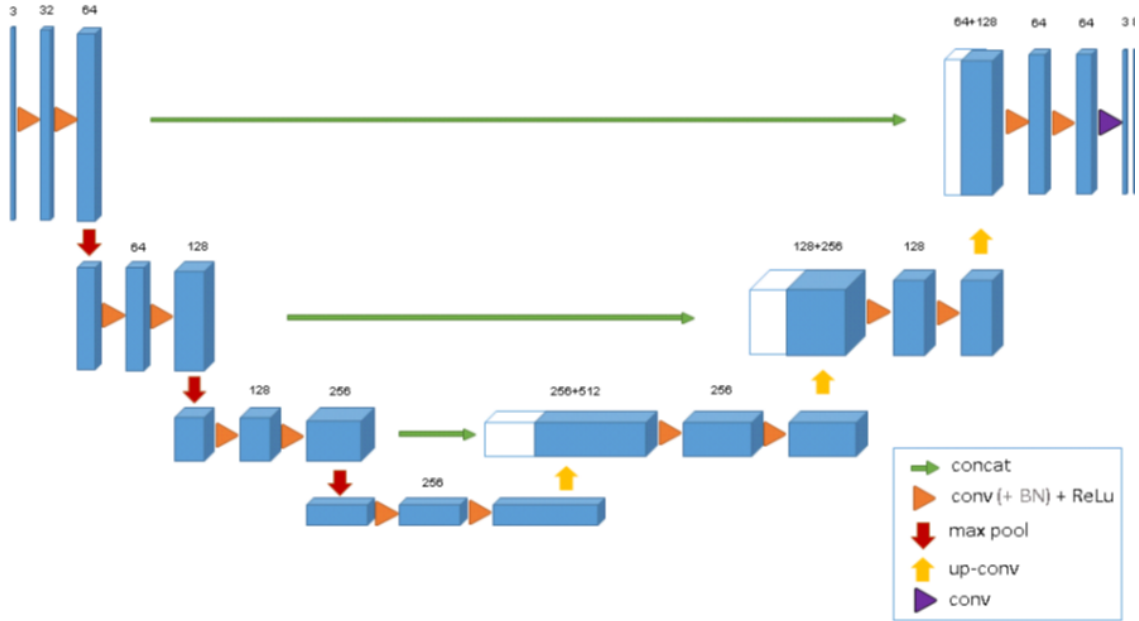


Fig. 3.4: The 3D U-Net network architecture [5].

This means one medical image can be labelled in the target domains to realise semantic segmentation. In this project, the framework is modified to fit the dataset magnitude.

1. In the analysis path, all four "Down" layers contain one  $2 \times 2 \times 2$  convolution layer alone with a stride of two in all dimensions, and three same convolution layers each followed by an activation function (ReLU or Softplus) and a normalisation layer (Instance Normalisation or Batch Normalisation). The max pooling layer in the original 3D U-Net model is not applied, which is replaced by an optional dropout layer.
2. In the synthesis path, all four "Up" layers consist of the up-convolution layers whose structure and parameters are similar to the "Down" ones but in the opposite direction.
3. Shortcut connections from layers of equal resolution between analysis and synthesis paths remain to import high-resolution features realised by concatenation.
4. In the last layer, a convolution layer reduces the number of output channels to the number of labels. Then, a Softmax function is attached to get the final labels for each point.

### 3.3.2 UNet Transformer

UNet Transformer (UNETR) follows 3D U-Net "U-shaped" network design for the encoder and decoder, but replaces the CNN layers with a ViT transformer in the encoder to consider volume sequence representations and capture the multi-dimension information [6].

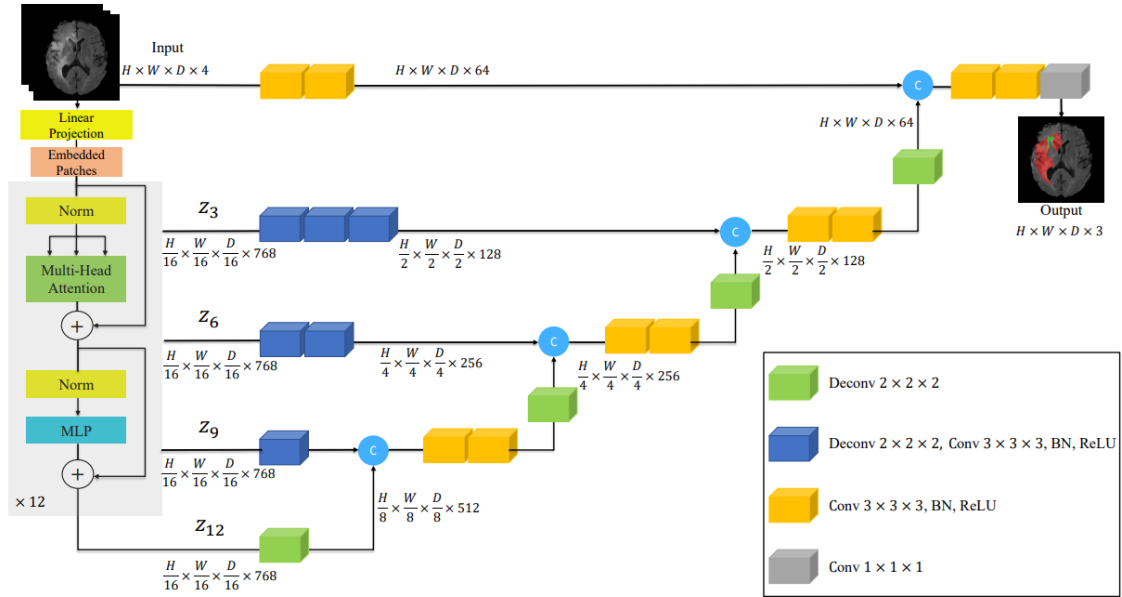


Fig. 3.5: The UNETR network architecture [6].

According to ViT, a 1D sequence of a 3D input volume is initialized by splitting it into flattened patches and embedded with its positional sequence at beginning [6]. In encoder path, the twelve transformer blocks are divided into four stack of transformers, which can be seen as encoder layers [6]. Similar to 3D U-Net, features from multiple resolutions of the encoder are merged with the decoder by concatenating volumes in the same level [6]. However, the hidden layer outputs

need to reshaped into the target volume size and encoded through some deconvolution blocks [6]. Compared with standard 3D U-Net, UNETR discards max-pooling layers between levels, which increases the parameter size.

## 3.4 Model Shaping Network

### 3.4.1 Variational Autoencoder

The Variational Autoencoder (VAE) is applied for the recognition, denoising, representation and visualisation by learning the approximate posterior inference model [7]. The VAE network consists of Encoder and Decoder at the architecture level [7]. To allow back propagation, the reparameterisation trick is introduced in the sampling function, which combines the distribution parameters (mean and standard deviation) and a random sampled number in a scale [7]. Through KL divergence, Gaussian distribution, and Bayes formula, the final result formula of the VAE can be presented as the one in Equation 3.5, under the reparameterisation trick [7]. This means one image can be sampled into features as a latent variable and re-sampled to restore the original image.

$$\begin{aligned} \mathcal{L}(\theta, \phi; x^{(i)}) &\simeq \frac{1}{2} \sum_{j=1}^J \left( 1 + \log \left( \left( \sigma_j^{(i)} \right)^2 \right) - \left( \mu_j^{(i)} \right)^2 - \left( \sigma_j^{(i)} \right)^2 \right) \\ &\quad + \frac{1}{L} \sum_{l=1}^L \log p_{\theta} \left( x^{(i)} \mid \mathbf{z}^{(i,l)} \right) \\ \text{where } \mathbf{z}^{(i,l)} &= \mu^{(i)} + \sigma^{(i)} \odot \epsilon^{(l)} \text{ and } \epsilon^{(l)} \sim \mathcal{N}(0, \mathbf{I}) \end{aligned} \quad (3.5)$$

In this project, the reconstructed mask, i.e., the shape information of labels, must be extracted by the VAE network trained with ground truth masks. The structure follows the original design of the VAE network but imports the Dice coefficient as the loss term.

1. In the encoder block, all five encoding layers contain one  $2 \times 2 \times 2$  convolution layer alone and three same convolution layers, followed by an activation function and a normalisation layer. The encoding layers and their parameters are analogous to the "Down" layer in the 3D U-Net, which reuse the layer-building function.
2. In the decode block, all five decoding layers consist of the up-convolution layers whose structure and parameters are similar to the encoding ones but in the opposite direction, which is analogous to the "Up" layer.
3. The mean and standard deviation are calculated in the middle layers for back propagation. For the reparameterisation trick, the random sample between zero and a given scale is optional.
4. In the last layer, a convolution layer reduces the number of output channels to the number of labels. Then, a Softmax function is attached to get the final labels for each point.

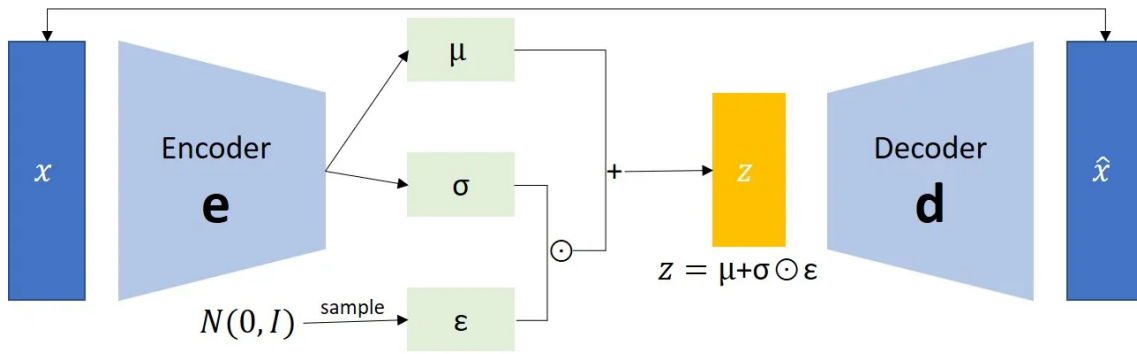


Fig. 3.6: The VAE architecture [7].

### 3.5 Loss Function

Tumour areas in this lung tumour segmentation task represent a very small fraction of the full image in some slices. Such unbalanced problems can cause improper fitting such as a tendency to mark all to non-tumour leading to false negative results, applying inappropriate loss functions [30]. This project prefers to apply the Dice loss for evaluation and validation.

#### 3.5.1 Dice Loss

The Dice score coefficient is a measure of overlap widely used to assess segmentation performance when ground truth is available. The Dice loss can be expressed as

$$DL_s = 1 - \frac{2|X \cap Y| + \epsilon}{|X| + |Y| + \epsilon} \quad (3.6)$$

where  $|X \cap Y|$  is the intersection between  $X$  and  $Y$ ,  $|X|$  and  $|Y|$  are the numbers of elements of  $X$  and  $Y$ , and  $\epsilon$  is the smoothing parameter for Laplace smoothing [30].

Multiple labels Dice loss for Tensor is implemented, as it is not realized in PyTorch.

```

1 def dice_coef_multiclass(inputs, targets, num_class, smooth=1):
2     dice = 0
3     for index in range(num_class):
4         dice += dice_coef(inputs[:, index, ...], targets[:, index, ...],
5                             smooth)
6     return 1 - dice / num_class
7
8 def dice_coef(inputs, targets, smooth=1):
9     inputs = inputs.flatten()
10    targets = targets.flatten()
11    intersection = (inputs * targets).sum()
12    dice = (2. * intersection + smooth) / (inputs.sum() + targets.sum()
13        + smooth)
14    return dice

```



## Chapter 4

# Experiments

### 4.1 Experiment Process

#### 4.1.1 Train on source domain

Three datasets (MSD, RIDER, and NSCLC) have been trained on the two segmentation models (3D U-Net and UNETR) and shaping model (VAE) respectively with relevant labelled segmentation masks to access the upper bound scores. The pre-trained models are saved for best model selection and further domain adaptation in target domain.

#### 4.1.2 Train on target domain

The pre-trained segmentation and shaping models of one dataset are selected as the source dataset to initialize parameters in the two fixed teacher networks. The other two datasets left are target datasets to fine-tune the student segmentation network, where their ground truth masks are neglected as they are in unsupervised domain adaptation. The three datasets are determined to be the source dataset to observe their performance.

### 4.2 Results

#### 4.2.1 Model Performance

Table 4.1 and Table 4.2 present the segmentation results on 3D U-Net and UNETR as their segmentation models, which are evaluated with mean Dice score. In detail, the data in columns have same target datasets. The upper bound score means the result in the segmentation model trained by relevant labelled datasets, which can be seen as ceiling score. The direct score means validate the dataset on the model trained by another dataset without any finetune. The UDA score is the final score through the complete model. Domain adaptation do polish up the score from direct validation. Moreover, due to more fitting on a larger dataset, the UDA even have better score on the small NSCLC dataset.

		MSD	RIDER	NSCLC
Upper Bound		0.6676	0.7879	0.6963
Direct	MSD	/	0.7247	0.7248
	RIDER	0.6537	/	0.7511
	NSCLC	0.6065	0.6382	/
UDA	MSD	/	<b>0.7415</b>	0.7145
	RIDER	<b>0.6625</b>	/	<b>0.7637</b>
	NSCLC	0.6065	0.6669	/

Table 4.1: Performance of this UDA segmentation method on 3D U-Net.

		MSD	RIDER	NSCLC
Upper Bound		0.6708	0.7568	0.7020
Direct	MSD	/	0.6811	0.7265
	RIDER	0.6547	/	0.7424
	NSCLC	0.5994	0.6192	/
UDA	MSD	/	<b>0.7326</b>	0.7386
	RIDER	<b>0.6624</b>	/	<b>0.7432</b>
	NSCLC	0.6041	0.6357	/

Table 4.2: Performance of this UDA segmentation method on UNETR.

### 4.2.2 Training Process

Data visualization on training process of validation score is shown in the Fig 4.1, Fig 4.2, and Fig 4.3, which are based on UNETR model. As the figures show, the models in source domain can converge quickly in 20 epochs and be stabilised to a upper limit with some fluctuations. The target domain do not provide reasonable finetune towards target datasets. All charts including training losses and validation scores on 3D U-Net and UNETR are listed in the Appendix A.

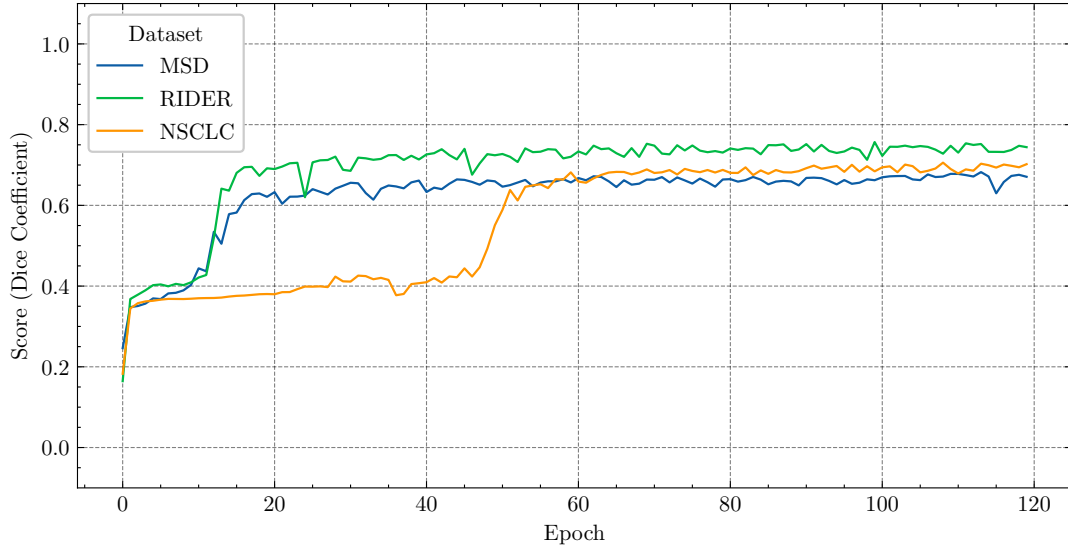


Fig. 4.1: The validation score chart for three datasets in UNETR segmentation model.

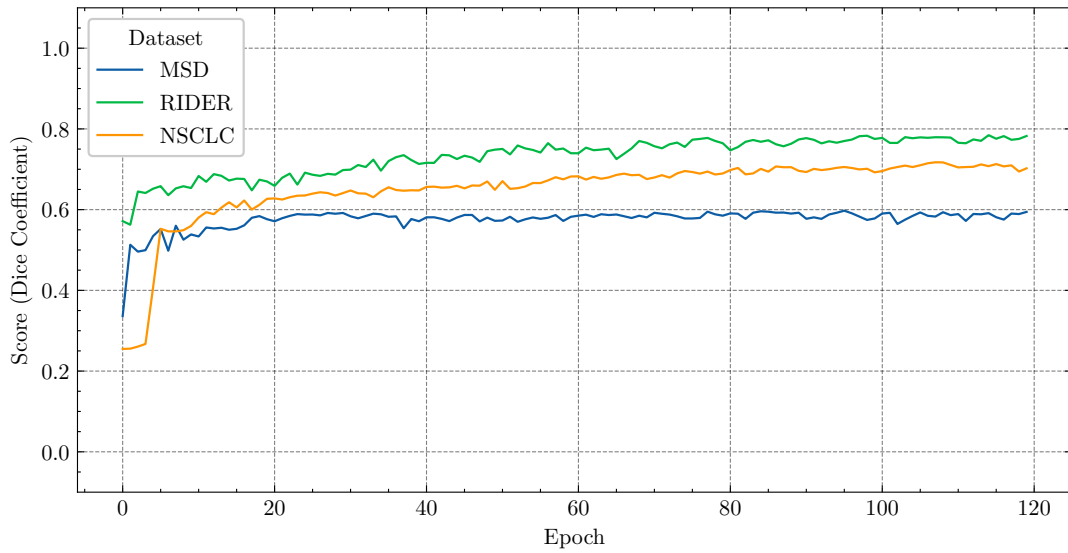


Fig. 4.2: The validation score chart for three datasets in VAE shaping model.

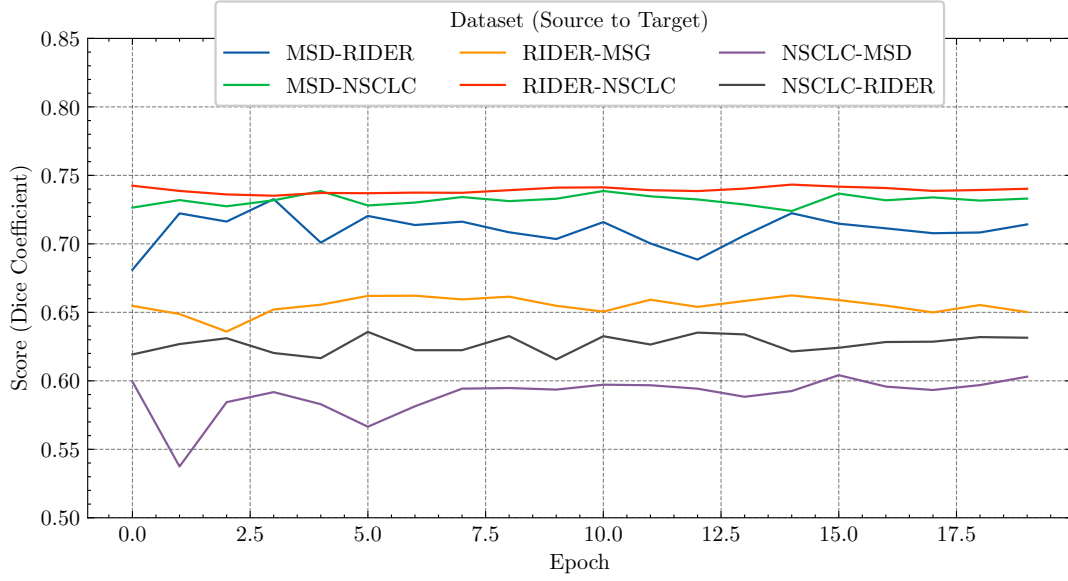


Fig. 4.3: The validation score chart for six dataset pairs in UDA framework based on UNETR.

#### 4.2.3 Visualization Result

As the original medical CT volumes are through data preprocessing and augmentation, it is hard for the outputs of the model to be converted to a standard medical segmentation format (i.e. NIfTI or DCMQI) for further 3D model reconstruction. Fig 4.4 and Fig 4.5 show two slices with different states for model performance display in both segmentation model and UDA framework.

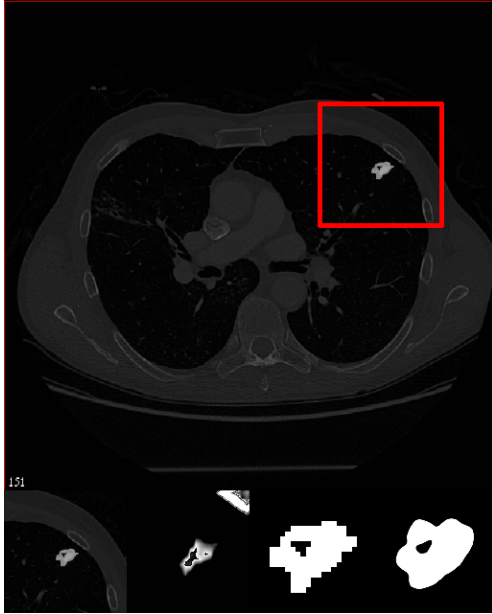


Fig. 4.4: The original segmentation result of "lung\_070.nii.gz" in the MSD dataset at frame 151 (from left to right, they are origin, preprocessed, label, and prediction).

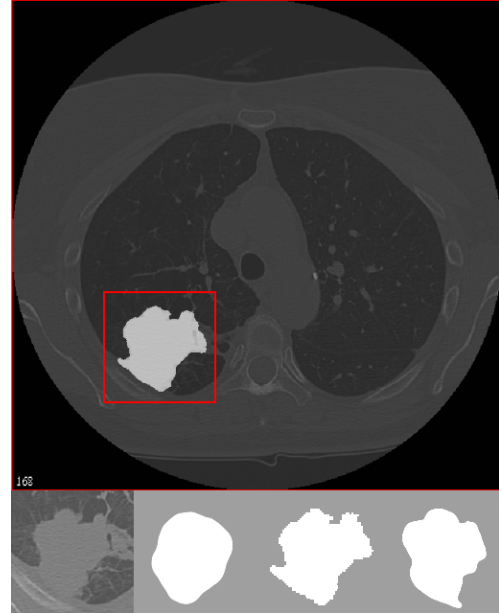


Fig. 4.5: The domain adaptation segmentation result of "RIDER-1129164940" in the RIDER dataset at frame 168 (from left to right, they are origin, reconstruction, label, and prediction).

## Chapter 5

# Conclusion and Future Work

### 5.1 Conclusion

In conclusion, the framework based on segmentation and shaping models can improve the model performance on unlabelled datasets by unsupervised domain adaptation. The Dice gap between a certain method and the upper bound is around 2% in most cases. The 3D U-Net with simple network architecture trained on the relatively abundant datasets RIDER reaches the highest scores among the current results, which satisfies the weakness of ViT in labelled dataset shortage. In the small dataset NSCLC, the model pre-trained on other datasets provide more accurate segmentation than the upper bound result trained by the same dataset.

### 5.2 Future Work

This project needs further modification and maintenance in the future. More datasets with different scanning styles, such as combination of CT and MRI, will be collected to analyse the bottlenecks of the current network. Some state-of-the-art components and mechanisms will be introduced into the network to handle the possible drawbacks, including accuracy, training time, and model size. Statistics about comparison with other segmentation methods will be evaluated in the future work.

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# Appendix A

## Charts

### A.1 3D U-Net Segmentation

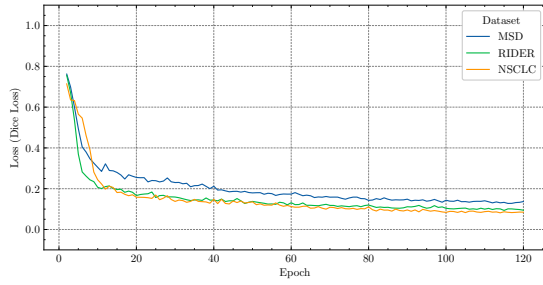


Fig. A.1: The training loss chart for three datasets in 3D U-Net segmentation model.

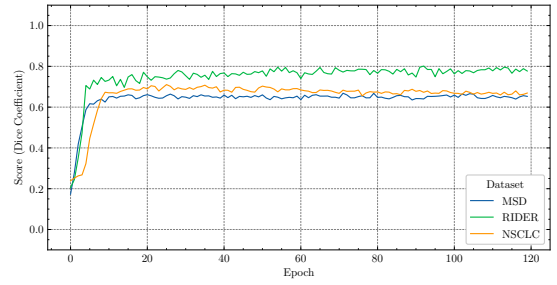


Fig. A.2: The validation score chart for three datasets in 3D U-Net segmentation model.

### A.2 UNETR Segmentation

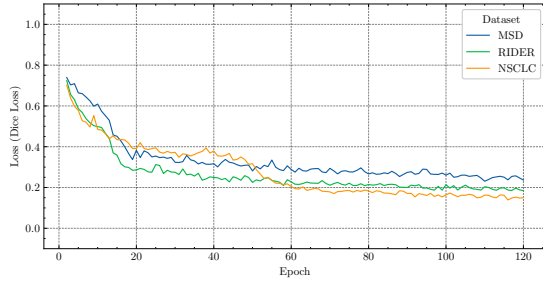


Fig. A.3: The training loss chart for three datasets in UNETR segmentation model.

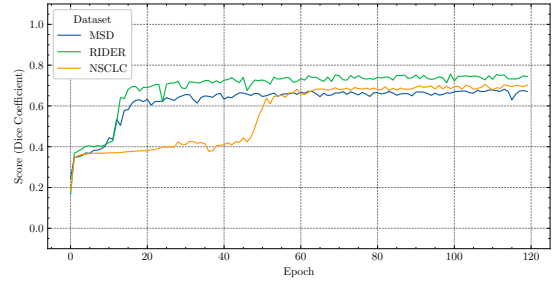


Fig. A.4: The validation score chart for three datasets in UNETR segmentation model.

### A.3 VAE Shaping

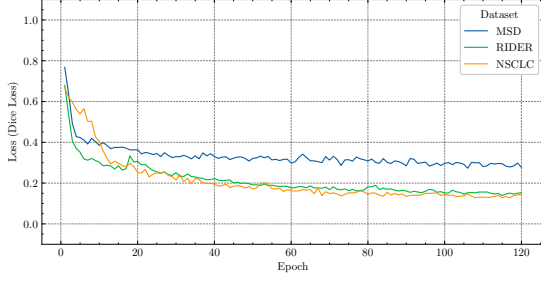


Fig. A.5: The training loss chart for three datasets in VAE shaping model.

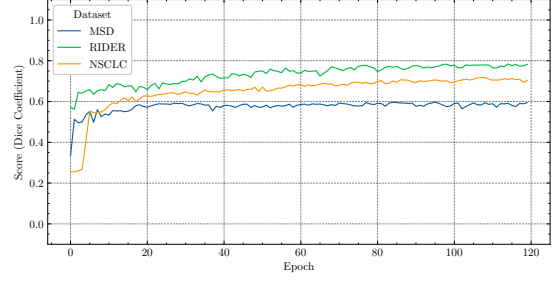


Fig. A.6: The validation score chart for three datasets in VAE shaping model.

### A.4 3D U-Net UDA framework

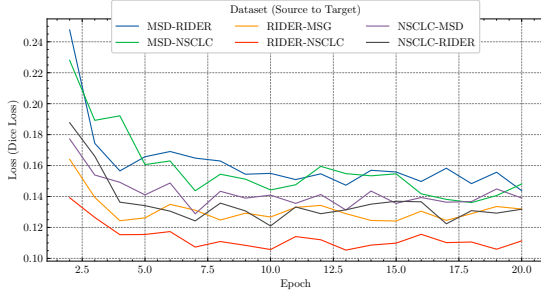


Fig. A.7: The training loss chart for six dataset pairs in 3D U-Net UDA framework.

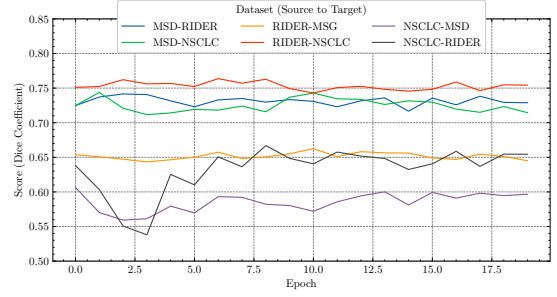


Fig. A.8: The validation score chart for six dataset pairs in 3D U-Net UDA framework.

### A.5 UNETR UDA framework

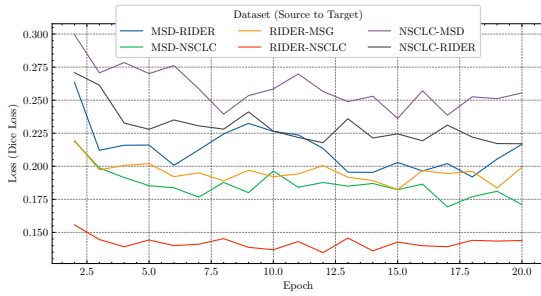


Fig. A.9: The training loss chart for six dataset pairs in UNETR UDA framework.

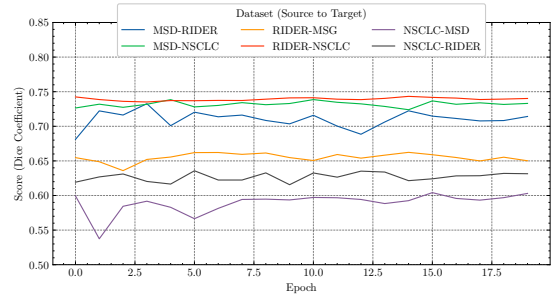


Fig. A.10: The validation score chart for six dataset pairs in UNETR UDA framework.

## Appendix B

# Project Code

### B.1 dcm\_to\_nii.py

```
1 import os
2
3 dcm2niix_path = ".\\tools\\MRICron\\Resources\\dcm2niix.exe"
4 dcm2niix_command = "{} -o "{}\\{}" -b n -z y -f "{}\\{}" "{}\\{}"
5
6 dcmqi_path = ".\\tools\\dcmqi\\bin\\segimage2itkimage.exe"
7 dcmqi_command = "{} --inputDICOM "{}\\{}" --outputDirectory "{}\\{}"
8     --prefix "{}\\{}" -t nifti"
9
10 ct_target_path = "./dataset/dcm_data/NSCLC1/CT"
11 ct_output_path = "./dataset/nih_data/NSCLC1/CT"
12 seg_target_path = "./dataset/dcm_data/NSCLC1/Seg"
13 seg_output_path = "./dataset/nih_data/NSCLC1/Seg"
14
15
16 def convert_ct():
17     for root, dirs, _ in os.walk(ct_target_path, topdown=False):
18         for name in dirs:
19             os.makedirs(ct_output_path, exist_ok=True)
20             os.system(dcm2niix_command.format(
21                 dcm2niix_path,
22                 ct_output_path,
23                 name,
24                 os.path.join(root, name),
25             ))
26
27
28 def convert_seg():
29     for root, dirs, files in os.walk(seg_target_path, topdown=False):
30         for name in files:
31             os.makedirs(seg_output_path, exist_ok=True)
32             os.system(dcmqi_command.format(
33                 dcmqi_path,
34                 os.path.join(root, name),
35                 seg_output_path,
```

```
36         os.path.basename(root)
37     ))
38
39
40 def main():
41     convert_ct()
42     convert_seg()
43
44
45 if __name__ == "__main__":
46     main()
```

## B.2 data\_process.py

```
1 import numpy as np
2 import nibabel as nib
3 import skimage
4 import os
5 import glob
6 import tqdm
7 import re
8
9 pad = [32, 32, 32]
10
11
12 def resample(nih, **kwargs):
13     scale = np.diagonal(nih.affine)[: 3]
14     image = np.flip(nih.get_fdata(), np.flatnonzero(scale > 0))
15     new_shape = (image.shape * np.abs(scale)).astype(int)
16     image = skimage.transform.resize(image.astype(np.float64),
17                                     new_shape, **kwargs)
18     return image
19
20 def get_box_index(label):
21     mask_index = np.array((label > 0).nonzero())
22     max_index = (mask_index.max(axis=1) + pad).clip(max=label.shape)
23     min_index = (mask_index.min(axis=1) - pad).clip(min=0)
24     center = np.ceil((max_index + min_index) / 2).astype(int)
25     length = np.ceil((max_index - min_index) / 2).max().astype(int)
26     max_index = (center + length).clip(max=label.shape)
27     min_index = (center - length).clip(min=0)
28     return max_index, min_index
29
30
31 def get_image_box(image, box):
32     max_index, min_index = box
33     return image[
34         min_index[0]:max_index[0], \
35         min_index[1]:max_index[1], \
36         min_index[2]:max_index[2], \
37     ]
38
39
40 def main():
41     image_path = '../dataset/nih_data/NSCLC1/CT'
42     label_path = '../dataset/nih_data/NSCLC1/Seg'
43     to_path = './nih/nsclcl_test'
44     os.makedirs(to_path, exist_ok=True)
45
46     names = glob.glob(os.path.join(image_path, '*.gz'))
47     names = [os.path.split(f)[1] for f in names]
48
49     for img_name in tqdm.tqdm(names):
50         label_name = re.sub(r"(?=.nii.gz)", "-1", img_name)
51
52         image = nib.load(os.path.join(image_path, img_name))
```

```

53     label = nib.load(os.path.join(label_path, label_name))
54
55     image = resample(image)
56     label = resample(label, anti_aliasing=False, order=0)
57
58     label_box = get_box_index(label)
59
60     image = get_image_box(image, label_box)
61     label = get_image_box(label, label_box)
62
63     path_prefix = os.path.join(to_path, img_name.split('.')[0])
64     os.makedirs(path_prefix, exist_ok=True)
65
66     np.save(os.path.join(path_prefix, 'img.npy'),
67             image.astype(np.int16))
68     np.save(os.path.join(path_prefix, 'label.npy'),
69             label.astype(np.int8))
70     np.save(os.path.join(path_prefix, 'merge.npy'),
71             np.stack((image, label), axis=-1).astype(np.int16))
72
73 if __name__ == '__main__':
74     main()

```

## B.3 main\_source.py

```
1 import argparse
2 import os
3
4 from batchgenerators.transforms.sample_normalization_transforms import
    RangeTransform
5 from batchgenerators.transforms.spatial_transforms import
    SpatialTransform
6 from batchgenerators.transforms.abstract_transforms import Compose
7 from monai.networks.nets import UNETR
8 import torch
9 from torch.utils.data import DataLoader
10 import tqdm
11
12 from models import Segmentation, VAE
13 from utils import CropResize, TensorBoardWriter, filedict_from_json,
    DiceLoss, NiiDataset, KL_loss
14
15
16 parser = argparse.ArgumentParser()
17 parser.add_argument("prefix")
18 parser.add_argument("--method", default='UNET_train')
19 parser.add_argument("--batch_size", type=int, default=2)
20 parser.add_argument("--max_epoch", type=int, default=120)
21 parser.add_argument("--save_epoch", type=int, default=20)
22 parser.add_argument("--data_index", default='./data/data_index.json')
23 parser.add_argument("--train_list", default='MSD_train')
24 parser.add_argument("--val_list", default='MSD_val')
25 parser.add_argument("--train_data_root", default='./data/nih/msd')
26 parser.add_argument("--val_data_root", default='./data/nih/msd')
27 parser.add_argument("--save_root", default='./model')
28 parser.add_argument("--display_root", default='./tensorboard')
29 parser.add_argument("--checkpoint_name", default="best_model.ckpt")
30 parser.add_argument("--lr", type=float, default=1e-2)
31 parser.add_argument("--weight_decay", type=float, default=0)
32 parser.add_argument("--test_only", action='store_true')
33 args = parser.parse_args()
34
35 prefix = args.prefix
36 method = args.method
37 train_batch = args.batch_size
38 val_batch = 1
39 max_epoch = args.max_epoch
40 save_epoch = args.save_epoch
41 data_index = args.data_index
42 train_list = args.train_list
43 val_list = args.val_list
44 train_data_root = args.train_data_root
45 val_data_root = args.val_data_root
46 save_path = os.path.join(args.save_root, prefix)
47 display_path = os.path.join(args.display_root, prefix)
48 checkpoint_name = args.checkpoint_name
49 lr = args.lr
50 weight_decay = args.weight_decay
```

```

51 test_only = args.test_only
52
53 num_workers = 4
54 torch.backends.cudnn.benchmark = True
55
56 patch_size = [128, 128, 128]
57 num_class = 2
58
59 for path in [save_path, display_path]:
60     os.makedirs(path, exist_ok=True)
61
62
63 def main():
64     print("Loading data")
65     train_data_list = filedict_from_json(data_index, train_list)
66     val_data_list = filedict_from_json(data_index, val_list)
67
68     transforms = {
69         "train": Compose([
70             CropResize(
71                 data_key="data",
72                 label_key="label",
73                 output_size=patch_size
74             ),
75             RangeTransform(
76                 data_key="data",
77                 label_key="label"
78             ),
79             SpatialTransform(
80                 patch_size,
81                 [dis // 2 - 5 for dis in patch_size],
82                 random_crop=True,
83                 scale=(0.85, 1.15),
84                 do_rotation=True,
85                 angle_x=(-0.2, 0.2),
86                 angle_y=(-0.2, 0.2),
87                 angle_z=(-0.2, 0.2),
88                 data_key="data",
89                 label_key="label",
90             ),
91         ]),
92         "val": Compose([
93             CropResize(
94                 data_key="data",
95                 label_key="label",
96                 output_size=patch_size
97             ),
98             RangeTransform(
99                 data_key="data",
100                 label_key="label"
101             ),
102             SpatialTransform(
103                 patch_size,
104                 do_rotation=False,
105                 do_scale=False,

```



```

106         do_elastic_deform=False,
107         random_crop=False,
108         data_key="data",
109         label_key="label"
110     ),
111 ])
112 }
113
114 train_dataset = NiiDataset(train_data_root, train_data_list,
115                             transforms["train"])
116 val_dataset = NiiDataset(val_data_root, val_data_list,
117                           transforms["val"])
118
119 train_loader = DataLoader(train_dataset, batch_size=train_batch,
120                           shuffle=True,
121                           num_workers=num_workers, drop_last=True,
122                           pin_memory=True)
123 val_loader = DataLoader(val_dataset, batch_size=val_batch,
124                         shuffle=False,
125                         num_workers=num_workers, pin_memory=True)
126
127 print("Building model")
128
129 if method == 'vae_train':
130     model = VAE(n_channels=1, n_class=num_class, norm_type=1,
131                dim=128).cuda()
132 elif method == 'unet_train':
133     model = Segmentation(n_channels=1, n_class=num_class,
134                          norm_type=1).cuda()
135 elif method == 'unetr_train':
136     model = UNETR(in_channels=1, out_channels=num_class,
137                   img_size=(128, 128, 128)).cuda()
138 else:
139     raise ValueError("Try a valid method")
140
141 criterion = DiceLoss(num_class=num_class)
142 optimizer = torch.optim.SGD(
143     model.parameters(),
144     lr=lr,
145     weight_decay=weight_decay,
146     momentum=0.9
147 )
148
149 best_result = 0
150 saver = TensorBoardWriter(display_path)
151
152 print("Start training")
153
154 for epoch in tqdm.tqdm(range(max_epoch)):
155     if not test_only:
156         model.train()
157         total_loss = 0.0
158         for idx, batch in enumerate(train_loader):
159             optimizer.zero_grad()

```

```

153     batch["data"] = batch["data"].cuda()
154     batch["label"] = batch["label"].cuda()
155     batch["label"] = batch["label"].type(torch.cuda.LongTensor)
156     one_hot = torch.cuda.FloatTensor(
157         batch["label"].size(0),
158         num_class,
159         batch["label"].size(2),
160         batch["label"].size(3),
161         batch["label"].size(4)
162     ).zero_()
163     batch["label"] = one_hot.scatter_(1, batch["label"].data,
164         1)
165
166     if method in ['unet_train', 'unetr_train']:
167         predict_mask = model(batch["data"])
168
169         dice_loss = criterion(batch["label"], predict_mask)
170         final_loss = dice_loss
171
172         print('%3d, %3d] loss: %.4f' %
173             (epoch + 1, idx + 1, final_loss.item()))
174     elif method in ['vae_train']:
175         reconstruct_mask, mean, std = model(batch["label"])
176
177         dice_loss = criterion(batch["label"], reconstruct_mask)
178         kl_loss = KL_loss(mean, std)
179         final_loss = dice_loss + 0.00002 * kl_loss
180
181         print('%3d, %3d] loss: %.4f, %.4f' %
182             (epoch + 1, idx + 1, dice_loss.item(),
183              kl_loss.item()))
184     else:
185         raise ValueError("Try a valid method")
186
187     total_loss += final_loss.item()
188     final_loss.backward()
189     optimizer.step()
190     saver.add_scale("train_loss", total_loss, epoch)
191
192     print("Start validation")
193
194     model.eval()
195     current_result = 0.0
196     with torch.no_grad():
197         for idx, batch in enumerate(val_loader):
198             batch["data"] = batch["data"].cuda()
199             batch['label'] = batch['label'].type(torch.cuda.LongTensor)
200             one_hot = torch.cuda.FloatTensor(
201                 batch['label'].size(0),
202                 num_class,
203                 batch['label'].size(2),
204                 batch['label'].size(3),
205                 batch['label'].size(4)
206             ).zero_()

```

```

205         batch['label'] = one_hot.scatter_(1, batch['label'].data,
206                                           1)
207
208         if method in ['unet_train', 'unetr_train']:
209             prediction = model(batch["data"])
210             current_result += criterion(batch["label"],
211                                       prediction).item()
212         elif method in ['vae_train']:
213             prediction, mean, std = model(batch["label"])
214
215             dice_loss = criterion(batch["label"], prediction)
216             kl_loss = KL_loss(mean, std)
217             current_result += (dice_loss + 0.00002 * kl_loss).item()
218         else:
219             raise ValueError("Try a valid method")
220
221         h = prediction.shape[4] // 2
222         saver.add_image(
223             "val_display",
224             torch.cat(
225                 (
226                     batch["data"][0:1, 0:1, :, :, h],
227                     batch["label"][0:1, 1:2, :, :, h],
228                     prediction[0:1, 1:2, :, :, h]
229                 ),
230                 dim=0
231             ),
232             idx + epoch * (len(batch))
233         )
234
235         current_result = 1 - current_result / (len(val_loader))
236
237         saver.add_scale("val_score", current_result, epoch)
238         print('epoch %d validation result: %f, best result %f.' %
239               (epoch + 1, current_result, best_result))
240
241     if test_only:
242         break
243
244     if (epoch + 1) % save_epoch == 0:
245         print('Saving model')
246         torch.save(
247             {
248                 'epoch': epoch + 1,
249                 'model_state_dict': model.state_dict(),
250                 'optimizer_state_dict': optimizer.state_dict()
251             },
252             os.path.join(save_path, f'model_epoch{epoch + 1}.ckpt')
253         )
254         if current_result > best_result:
255             best_result = current_result
256             torch.save(
257                 {
258                     'epoch': epoch + 1,
259                     'model_state_dict': model.state_dict(),

```

```
258         'optimizer_state_dict': optimizer.state_dict()
259     },
260     os.path.join(save_path, 'best_model.ckpt')
261 )
262
263 print('Finished Training')
264
265
266 if __name__ == "__main__":
267     main()
```

## B.4 main\_target.py

```
1 import argparse
2 import os
3
4 from batchgenerators.transforms.sample_normalization_transforms import
    RangeTransform
5 from batchgenerators.transforms.spatial_transforms import
    SpatialTransform
6 from batchgenerators.transforms.abstract_transforms import Compose
7 from monai.networks.nets import UNETR
8 import torch
9 from torch.utils.data import DataLoader
10 import tqdm
11
12 from models import Segmentation, VAE
13 from utils import NiiDataset, filedict_from_json, CropResize,
    TensorBoardWriter, DiceLoss
14
15
16 parser = argparse.ArgumentParser()
17 parser.add_argument("prefix")
18 parser.add_argument("--method", default='UNET_train')
19 parser.add_argument("--batch_size", type=int, default=2)
20 parser.add_argument("--max_epoch", type=int, default=120)
21 parser.add_argument("--save_epoch", type=int, default=20)
22 parser.add_argument("--data_index", default='./data/data_index.json')
23 parser.add_argument("--train_list", default='MSD_train')
24 parser.add_argument("--val_list", default='MSD_val')
25 parser.add_argument("--train_data_root", default='./data/nih/msd')
26 parser.add_argument("--val_data_root", default='./data/nih/msd')
27 parser.add_argument("--save_root", default='./model')
28 parser.add_argument("--display_root", default='./tensorboard')
29 parser.add_argument("--checkpoint_name", default="best_model.ckpt")
30 parser.add_argument("--lr", type=float, default=1e-2)
31 parser.add_argument("--weight_decay", type=float, default=0)
32 parser.add_argument("--lambda_vae", type=float, default=0.1)
33 parser.add_argument("--test_only", action='store_true')
34 parser.add_argument("--load_prefix_seg", default=None)
35 parser.add_argument("--load_prefix_vae", default=None)
36 args = parser.parse_args()
37
38 prefix = args.prefix
39 method = args.method
40 train_batch = args.batch_size
41 val_batch = 1
42 max_epoch = args.max_epoch
43 save_epoch = args.save_epoch
44 data_index = args.data_index
45 train_list = args.train_list
46 val_list = args.val_list
47 train_data_root = args.train_data_root
48 val_data_root = args.val_data_root
49 save_root = args.save_root
50 save_path = os.path.join(save_root, prefix)
```

```

51 display_path = os.path.join(args.display_root, prefix)
52 checkpoint_name = args.checkpoint_name
53 lr = args.lr
54 weight_decay = args.weight_decay
55 lambda_vae = args.lambda_vae
56 test_only = args.test_only
57 load_prefix_seg = args.load_prefix_seg
58 load_prefix_vae = args.load_prefix_vae
59
60 num_workers = 4
61 torch.backends.cudnn.benchmark = True
62
63 patch_size = [128, 128, 128]
64 num_class = 2
65
66 for path in [save_path, display_path]:
67     os.makedirs(path, exist_ok=True)
68
69
70 def main():
71     train_data_list = filedict_from_json(data_index, train_list)
72     val_data_list = filedict_from_json(data_index, val_list)
73
74     transforms = {
75         "train": Compose([
76             CropResize(
77                 data_key="data",
78                 label_key="label",
79                 output_size=patch_size
80             ),
81             RangeTransform(
82                 data_key="data",
83                 label_key="label"
84             ),
85             SpatialTransform(
86                 patch_size,
87                 [dis // 2 - 5 for dis in patch_size],
88                 random_crop=True,
89                 scale=(0.85, 1.15),
90                 do_rotation=True,
91                 angle_x=(-0.2, 0.2),
92                 angle_y=(-0.2, 0.2),
93                 angle_z=(-0.2, 0.2),
94                 data_key="data",
95                 label_key="label",
96             ),
97         ]),
98         "val": Compose([
99             CropResize(
100                 data_key="data",
101                 label_key="label",
102                 output_size=patch_size
103             ),
104             RangeTransform(
105                 data_key="data",

```

```

106         label_key="label"
107     ),
108     SpatialTransform(
109         patch_size,
110         do_rotation=False,
111         do_scale=False,
112         do_elastic_deform=False,
113         random_crop=False,
114         data_key="data",
115         label_key="label"
116     ),
117 ])
118 }
119
120 print("Loading data")
121
122 train_dataset = NiiDataset(train_data_root, train_data_list,
123                             transforms["train"])
124 val_dataset = NiiDataset(val_data_root, val_data_list,
125                           transforms["val"])
126
127 train_loader = DataLoader(train_dataset, batch_size=train_batch,
128                           shuffle=True,
129                           num_workers=num_workers, drop_last=True,
130                           pin_memory=True)
131 val_loader = DataLoader(val_dataset, batch_size=val_batch,
132                         shuffle=False,
133                         num_workers=num_workers, pin_memory=True)
134
135 print("Building model")
136
137 if method in ["unet_train"]:
138     model = {
139         "teacher": {
140             "seg": Segmentation(n_channels=1, n_class=num_class,
141                                norm_type=1).cuda(),
142             "vae": VAE(n_channels=1, n_class=num_class, norm_type=1,
143                       dim=128).cuda(),
144         },
145         "student": {
146             "seg": Segmentation(n_channels=1, n_class=num_class,
147                                norm_type=1).cuda()
148         }
149     }
150 elif method in ["unetr_train"]:
151     model = {
152         "teacher": {
153             "seg": UNETR(in_channels=1, out_channels=num_class,
154                          img_size=(128, 128, 128)).cuda(),
155             "vae": VAE(n_channels=1, n_class=num_class, norm_type=1,
156                       dim=128).cuda(),
157         },
158         "student": {
159             "seg": UNETR(in_channels=1, out_channels=num_class,
160                          img_size=(128, 128, 128)).cuda()
161         }
162     }

```

```

150     }
151 }
152 else:
153     raise ValueError("Try a valid method")
154
155 criterion = DiceLoss(num_class=num_class)
156 optimizer = torch.optim.SGD(model["student"]["seg"].parameters(),
157                             lr=lr, weight_decay=weight_decay, momentum=0.9)
158
159 print("Loading prefix")
160
161 if load_prefix_seg:
162     model_path = os.path.join(save_root, load_prefix_seg,
163                               checkpoint_name)
164     model_state_dict = torch.load(model_path)['model_state_dict']
165     model["teacher"]["seg"].load_state_dict(model_state_dict)
166     model["student"]["seg"].load_state_dict(model_state_dict)
167
168 if load_prefix_vae:
169     model_path = os.path.join(save_root, load_prefix_vae,
170                               checkpoint_name)
171     model_state_dict = torch.load(model_path)['model_state_dict']
172     model["teacher"]["vae"].load_state_dict(model_state_dict)
173
174 for item in model["teacher"].values():
175     for param in item.parameters():
176         param.requires_grad = False
177     item.eval()
178
179 best_result = 0
180 saver = TensorBoardWriter(display_path)
181
182 print("Start training")
183
184 for epoch in tqdm.tqdm(range(max_epoch)):
185     if not test_only:
186         model["student"]["seg"].train()
187         total_final_loss = 0.0
188         for idx, batch in enumerate(train_loader):
189             optimizer.zero_grad()
190             model["student"]["seg"].train()
191             batch["data"] = batch["data"].cuda()
192             batch["label"] = batch["label"].cuda()
193             batch["label"] = batch["label"].type(torch.cuda.LongTensor)
194             one_hot = torch.cuda.FloatTensor(
195                 batch["label"].size(0),
196                 num_class,
197                 batch["label"].size(2),
198                 batch["label"].size(3),
199                 batch["label"].size(4)
200             ).zero_()
201             batch["label"] = one_hot.scatter_(1, batch["label"].data,
202                                             1)
203
204             pseudo_label = model["teacher"]["seg"](batch["data"])

```



```

202     predict_mask = model["student"]["seg"](batch["data"])
203     restruct_predict_mask, _, _ =
        model["teacher"]["vae"](predict_mask)
204
205     recon_loss = criterion(restruct_predict_mask, predict_mask)
206     pseudo_loss = criterion(pseudo_label, predict_mask)
207     final_loss = lambda_vae * recon_loss + pseudo_loss
208     total_final_loss += final_loss
209
210     print('[%3d, %3d] loss: %.4f, %.4f, %.4f' %
211           (epoch + 1, idx + 1, final_loss, recon_loss,
212            pseudo_loss))
213
214     final_loss.backward()
215     optimizer.step()
216
217     saver.add_scale("train_loss", total_final_loss, epoch)
218
219     print("Start validation")
220
221     model["student"]["seg"].eval()
222     current_result = 0.0
223     with torch.no_grad():
224         for idx, batch in enumerate(val_loader):
225             batch["data"] = batch["data"].cuda()
226             batch["label"] = batch["label"].cuda()
227             batch["label"] = batch["label"].type(torch.cuda.LongTensor)
228             one_hot = torch.cuda.FloatTensor(
229                 batch["label"].size(0),
230                 num_class,
231                 batch["label"].size(2),
232                 batch["label"].size(3),
233                 batch["label"].size(4)
234             ).zero_()
235             batch["label"] = one_hot.scatter_(1, batch["label"].data,
236                                             1)
237
238             pseudo_label = model["teacher"]["seg"](batch["data"])
239             predict_mask = model["student"]["seg"](batch["data"])
240             restruct_predict_mask, _, _ =
241                 model["teacher"]["vae"](predict_mask, if_random=False,
242                                         scale=0)
243
244             dice_loss = criterion(predict_mask, batch["label"])
245             current_result += dice_loss
246
247             h = predict_mask.shape[4] // 2
248             saver.add_image(
249                 "val_display",
250                 torch.cat(
251                     (
252                         batch["data"][0:1, 0:1, :, :, h],
253                         batch["label"][0:1, 1:2, :, :, h],
254                         predict_mask[0:1, 1:2, :, :, h],
255                         pseudo_label[0:1, 1:2, :, :, h],

```

```

252         reconstruct_predict_mask[0:1, 1:2, :, :, h],
253     ),
254     dim=0
255 ),
256     idx + epoch * (len(batch))
257 )
258
259     current_result = 1 - current_result / (len(val_loader))
260
261     saver.add_scale("val_score", current_result, epoch)
262     print('epoch %d validation result: %f, best result %f.' %
263           (epoch + 1, current_result, best_result))
264
265     if test_only:
266         break
267
268     if (epoch + 1) % save_epoch == 0:
269         print('Saving model')
270         torch.save(
271             {
272                 'epoch': epoch + 1,
273                 'model_state_dict':
274                     model["student"]["seg"].state_dict(),
275                 'optimizer_state_dict': optimizer.state_dict()
276             },
277             os.path.join(save_path, f'model_epoch{epoch + 1}.ckpt')
278         )
279         if current_result > best_result:
280             best_result = current_result
281             torch.save(
282                 {
283                     'epoch': epoch + 1,
284                     'model_state_dict':
285                         model["student"]["seg"].state_dict(),
286                     'optimizer_state_dict': optimizer.state_dict()
287                 },
288                 os.path.join(save_path, 'best_model.ckpt')
289             )
290
291         print('Finished Training')
292
293 if __name__ == "__main__":
294     main()

```

## B.5 models.py

```
1 import torch
2
3
4 def Normalization(norm_type, out_channels, num_group=1):
5     if norm_type == 1:
6         return torch.nn.InstanceNorm3d(out_channels)
7     elif norm_type == 2:
8         return torch.nn.BatchNorm3d(out_channels, momentum=0.1)
9
10
11 class DoubleConv(torch.nn.Module):
12     def __init__(self, in_ch, out_ch, norm_type=2, soft=False):
13         super().__init__()
14         activation = torch.nn.Softplus() if soft else
15             torch.nn.ReLU(inplace=False)
16         self.conv = torch.nn.Sequential(
17             torch.nn.Conv3d(in_ch, out_ch, 3, padding=1),
18             Normalization(norm_type, out_ch),
19             activation,
20             torch.nn.Conv3d(out_ch, out_ch, 3, padding=1),
21             Normalization(norm_type, out_ch),
22             activation,
23             torch.nn.Conv3d(out_ch, out_ch, 3, padding=1),
24             Normalization(norm_type, out_ch),
25             activation
26         )
27
28     def forward(self, x):
29         x = self.conv(x)
30         return x
31
32 class Conv(torch.nn.Module):
33     def __init__(self, in_ch, out_ch, norm_type=2, num_group=1,
34         activation=True, norm=True,
35         soft=False):
36         super().__init__()
37         activation = torch.nn.Softplus() if soft else
38             torch.nn.ReLU(inplace=True)
39         self.conv = torch.nn.Sequential(
40             torch.nn.Conv3d(in_ch, out_ch, 3, padding=1),
41             Normalization(norm_type, out_ch),
42             activation,
43         )
44
45     def forward(self, x):
46         x = self.conv(x)
47         return x
48
49 class Up(torch.nn.Module):
50     def __init__(self, in_ch, out_ch, norm_type=2, kernal_size=(2, 2,
51         2), stride=(2, 2, 2),
```

```

50         soft=False):
51     super().__init__()
52     self.conv = torch.nn.Sequential(
53         torch.nn.ConvTranspose3d(in_ch, in_ch, kernal_size,
54             stride=stride, padding=0),
55         DoubleConv(in_ch, out_ch, norm_type, soft=False)
56     )
57
58     def forward(self, x):
59         x = self.conv(x)
60         return x
61
62 class Down(torch.nn.Module):
63     def __init__(self, in_ch, out_ch, norm_type=2, kernal_size=(2, 2,
64         2), stride=(2, 2, 2),
65         soft=False):
66     super().__init__()
67     self.conv = torch.nn.Sequential(
68         torch.nn.Conv3d(in_ch, in_ch, kernal_size, stride=stride,
69             padding=0),
70         DoubleConv(in_ch, out_ch, norm_type, soft=False)
71     )
72
73     def forward(self, x):
74         x = self.conv(x)
75         return x
76
77 class VAE(torch.nn.Module):
78     def __init__(self, n_channels, n_class, norm_type=2, n_fmaps=None,
79         dim=1024, soft=False):
80     super().__init__()
81     if n_fmaps is None:
82         n_fmaps = [8, 16, 32, 64, 128, 256]
83     self.in_block = Conv(n_class, n_fmaps[0], norm_type=norm_type,
84         soft=False)
85     self.down1 = Down(n_fmaps[0], n_fmaps[1], norm_type=norm_type,
86         soft=False)
87     self.down2 = Down(n_fmaps[1], n_fmaps[2], norm_type=norm_type,
88         soft=False)
89     self.down3 = Down(n_fmaps[2], n_fmaps[3], norm_type=norm_type,
90         soft=False)
91     self.down4 = Down(n_fmaps[3], n_fmaps[4], norm_type=norm_type,
92         soft=False)
93     self.down5 = Down(n_fmaps[4], n_fmaps[5], norm_type=norm_type,
94         soft=False)
95     self.fc_mean = torch.nn.Linear(16384, dim)
96     self.fc_std = torch.nn.Linear(16384, dim)
97     self.fc2 = torch.nn.Linear(dim, 16384)
98     self.up1 = Up(n_fmaps[5], n_fmaps[4], norm_type=norm_type,
99         soft=False)
100    self.up2 = Up(n_fmaps[4], n_fmaps[3], norm_type=norm_type,
101        soft=False)

```

```

93     self.up3 = Up(n_fmaps[3], n_fmaps[2], norm_type=norm_type,
94                  soft=False)
95     self.up4 = Up(n_fmaps[2], n_fmaps[1], norm_type=norm_type,
96                  soft=False)
97     self.up5 = Up(n_fmaps[1], n_fmaps[0], norm_type=norm_type,
98                  soft=False)
99     self.out_block = torch.nn.Conv3d(n_fmaps[0], n_class, 3,
100                                     padding=1)
101     self.final = torch.nn.Softmax(dim=1)
102     self.n_class = n_class
103
104 def forward(self, x, if_random=False, scale=1, mid_input=False,
105            dropout=0.0):
106     if not mid_input:
107         x = self.in_block(x)
108         x = self.down1(x)
109         x = self.down2(x)
110         x = self.down3(x)
111         x = self.down4(x)
112         x = self.down5(x)
113         x = x.view(x.size(0), 16384)
114         x_mean = self.fc_mean(x)
115         x_std = torch.nn.ReLU()(self.fc_std(x))
116         z = torch.randn(x_mean.size(0),
117                        x_mean.size(1)).type(torch.cuda.FloatTensor)
118         if if_random:
119             x = self.fc2(x_mean + z * x_std * scale)
120         else:
121             x = self.fc2(x_mean)
122     else:
123         x = self.fc2(x)
124     x = x.view(x.size(0), 256, 4, 4, 4)
125
126     x = self.up1(x)
127     if dropout: x = torch.nn.functional.dropout(x, p=dropout,
128                                                training=True)
129     x = self.up2(x)
130     if dropout: x = torch.nn.functional.dropout(x, p=dropout,
131                                                training=True)
132     x = self.up3(x)
133     if dropout: x = torch.nn.functional.dropout(x, p=dropout,
134                                                training=True)
135     x = self.up4(x)
136     if dropout: x = torch.nn.functional.dropout(x, p=dropout,
137                                                training=True)
138     x = self.up5(x)
139     if dropout: x = torch.nn.functional.dropout(x, p=dropout,
140                                                training=True)
141     x = self.out_block(x)
142     x = self.final(x)
143
144     if not mid_input:
145         return x, x_mean, x_std
146     else:
147         return x

```

```

137
138
139 class Segmentation(torch.nn.Module):
140     def __init__(self, n_channels, n_class, norm_type=2, n_fmaps=None):
141         super().__init__()
142         if n_fmaps is None:
143             n_fmaps = [8, 16, 32, 64, 128, 256]
144             self.in_block = Conv(n_channels, n_fmaps[0],
145                                 norm_type=norm_type, soft=False)
146             self.down1 = Down(n_fmaps[0], n_fmaps[1], norm_type=norm_type,
147                               soft=False)
148             self.down2 = Down(n_fmaps[1], n_fmaps[2], norm_type=norm_type,
149                               soft=False)
150             self.down3 = Down(n_fmaps[2], n_fmaps[3], norm_type=norm_type,
151                               soft=False)
152             self.down4 = Down(n_fmaps[3], n_fmaps[4], norm_type=norm_type,
153                               soft=False)
154
155             self.up2 = Up(n_fmaps[4], n_fmaps[3], norm_type=norm_type,
156                           soft=False)
157             self.up3 = Up(n_fmaps[3], n_fmaps[2], norm_type=norm_type,
158                           soft=False)
159             self.up4 = Up(n_fmaps[2], n_fmaps[1], norm_type=norm_type,
160                           soft=False)
161             self.up5 = Up(n_fmaps[1], n_fmaps[0], norm_type=norm_type,
162                           soft=False)
163             self.out_block = torch.nn.Conv3d(n_fmaps[0], n_class, 3,
164                                                padding=1)
165             self.final = torch.nn.Softmax(dim=1)
166             self.n_class = n_class
167
168     def forward(self, x, dropout=0.0):
169         x1 = self.in_block(x)
170         x2 = self.down1(x1)
171         x3 = self.down2(x2)
172         x4 = self.down3(x3)
173         x5 = self.down4(x4)
174         x = self.up2(x5)
175         if dropout: x = torch.nn.functional.dropout(x, p=dropout,
176                                                       training=True)
177         x = self.up3(x) + x3
178         if dropout: x = torch.nn.functional.dropout(x, p=dropout,
179                                                       training=True)
180         x = self.up4(x) + x2
181         if dropout: x = torch.nn.functional.dropout(x, p=dropout,
182                                                       training=True)
183         x = self.up5(x)
184         if dropout: x = torch.nn.functional.dropout(x, p=dropout,
185                                                       training=True)
186         x = self.out_block(x)
187         if dropout: x = torch.nn.functional.dropout(x, p=dropout,
188                                                       training=True)
189         x = self.final(x)
190         return x

```

## B.6 utils.py

```
1 import json
2 import os
3
4 from batchgenerators.transforms.abstract_transforms import
    AbstractTransform
5 import numpy as np
6 from skimage.transform import resize
7 from tensorboardX import SummaryWriter
8 import torch
9 from torch import nn
10 from torch.utils.data import Dataset
11 import torchvision
12
13
14 def filedict_from_json(json_path, key):
15     with open(json_path, 'r') as f:
16         json_dict = json.load(f)
17         listdict = json_dict.get(key, [])
18     return listdict
19
20
21 class NiiDataset(Dataset):
22     def __init__(self, root_dir, img_dirs, transform=None):
23         self.root_dir = root_dir
24         self.img_dirs = img_dirs
25         self.transform = transform
26
27     def __len__(self):
28         return len(self.img_dirs)
29
30     def __getitem__(self, idx):
31         merge_data = np.load(os.path.join(self.root_dir,
32                                             self.img_dirs[idx]))
33         image = merge_data[np.newaxis, np.newaxis, ...,
34                             0].astype(np.float32)
35         label = merge_data[np.newaxis, np.newaxis, ...,
36                             1].astype(np.float32)
37         data = {"data": image, "label": label}
38         if self.transform:
39             data = self.transform(**data)
40         data = {"data": np.squeeze(data["data"], 0), "label":
41                 np.squeeze(data["label"], 0)}
42         return data
43
44
45 class DiceLoss(nn.Module):
46     def __init__(self, num_class, smooth=1):
47         super().__init__()
48         self.num_class = num_class
49         self.smooth = smooth
50
51     def forward(self, inputs, targets):
52         dice = 0
```

```

49     for index in range(self.num_class):
50         dice += self.dice_coef(inputs[:, index, ...], targets[:,
51                                index, ...])
52     return 1 - dice / self.num_class
53
54 def dice_coef(self, inputs, targets):
55     # flatten label and prediction tensors
56     inputs = inputs.flatten()
57     targets = targets.flatten()
58
59     intersection = (inputs * targets).sum()
60     dice = (2. * intersection + self.smooth) / (inputs.sum() +
61         targets.sum() + self.smooth)
62
63     return dice
64
65 class CropResize(AbstractTransform):
66     def __init__(self, output_size, data_key="data", label_key="label"):
67         self.output_size = output_size
68         self.data_key = data_key
69         self.label_key = label_key
70
71     def __call__(self, **data_dict):
72         image = np.squeeze(data_dict[self.data_key], axis=(0, 1))
73         label = np.squeeze(data_dict[self.label_key], axis=(0, 1))
74
75         label_box = self.get_box_index(label)
76
77         image = self.get_image_box(image, label_box)
78         label = self.get_image_box(label, label_box)
79
80         image = resize(image, self.output_size)
81         label = resize(label, self.output_size, order=0,
82                        anti_aliasing=False)
83
84         data_dict[self.data_key] = np.expand_dims(image, (0, 1))
85         data_dict[self.label_key] = np.expand_dims(label, (0, 1))
86
87         return data_dict
88
89     @staticmethod
90     def get_box_index(label):
91         mask_index = np.array((label > 0).nonzero())
92         max_index = (mask_index.max(axis=1)).clip(max=label.shape)
93         min_index = (mask_index.min(axis=1)).clip(min=0)
94         center = np.ceil((max_index + min_index) / 2).astype(int)
95         length = np.ceil((max_index - min_index) / 2).max().astype(int)
96         pad = int(length * 0.1)
97         max_index = (center + length + pad).clip(max=label.shape)
98         min_index = (center - length - pad).clip(min=0)
99         return max_index, min_index
100
101     @staticmethod
102     def get_image_box(image, box):

```



```

102     max_index, min_index = box
103     return image[
104         min_index[0]: max_index[0],
105         min_index[1]: max_index[1],
106         min_index[2]: max_index[2],
107     ]
108
109
110 def KL_loss(mean, std):
111     return torch.mean(0.5 * (
112         torch.sum(torch.pow(std, 2), 1)
113         + torch.sum(torch.pow(mean, 2), 1)
114         - 2 * torch.sum(torch.log(std + 0.00001), 1)
115     ))
116
117
118 class TensorBoardWriter():
119     def __init__(self, logdir):
120         self.writer = SummaryWriter(logdir=logdir)
121
122     def add_image(self, tag, image, step):
123         self.writer.add_image(
124             tag,
125             torchvision.utils.make_grid(image.detach()),
126             step
127         )
128
129     def add_scale(self, tag, num, step):
130         self.writer.add_scalar(
131             tag,
132             num,
133             step
134         )

```