

# Do not Treat Boundaries and Regions Differently: An Example on Heart Left Atrial Segmentation

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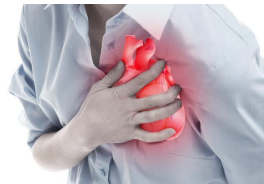
# Introduction

**Cardiovascular diseases** are the leading cause of death globally.

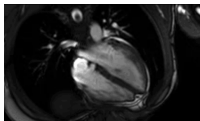
— — — according to the World Health Organization (WHO).

## Medical imaging

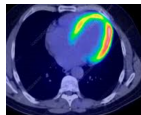
- Computed tomography (CT)
- Magnetic resonance imaging (MRI)
- Positron emission tomography (PET)
- Single photon emission computed tomography (SPECT)
- Ultrasound (US)



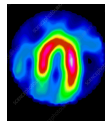
CT



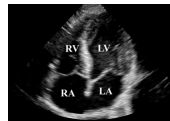
MRI



PET



SPECT



US

MRI: **Great contrast** between soft tissues and relatively high spatial resolutions.

## The difficulties of cardiac segmentation on the MR Images are shown as follow:

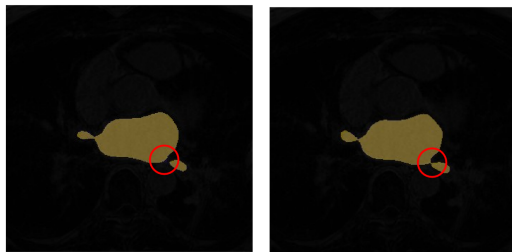
- **Poor contrast**
  - between myocardium and surrounding structures
- **Brightness**
  - in left ventricular/right ventricular cavities due to blood flow
- **Non-homogeneous partial volume**
  - limited CMR resolution(1.5T, 3.0T...) along the long-axis
- **Noise**
  - motion artifacts and heart dynamics
- **Shape and intensity variability**
  - different patients and pathologies





## Loss Function: Dice and Cross-entropy (CE)

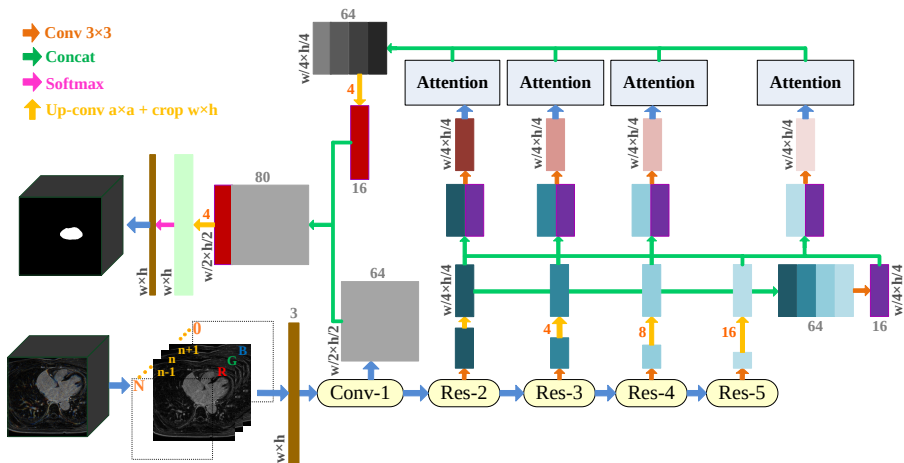
- CE: highly unbalanced problems.
- dice: difficulty dealing with very small structures.
- region-based.



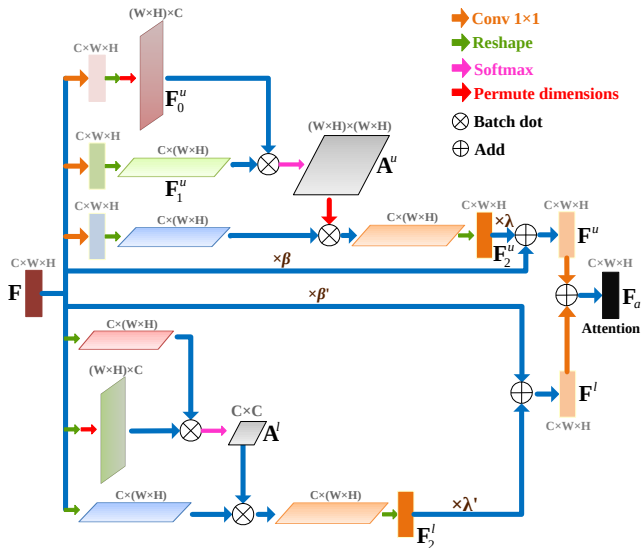
## There are three challenging problems applied on cardiac imaging:

- **enlarging** the receptive field of a CNN.
- **improving** the segmentation accuracy on small parts of objects.
- **balancing** the importance of the regions and the boundaries of objects.

# Methodology



Architecture of our networks.



**Attention Module.**  $\lambda$ ,  $\lambda'$ ,  $\beta$  and  $\beta'$  as hyperparameters, which is trained like the convolutional kernel. They decrease the weight of the unimportant feature maps.

# Hybrid Loss

The **hybrid loss** consists of two parts: **region loss** and **boundary loss**.

**Region Loss:**  $\ell_R = \ell_{\text{CCE}} + \ell_{\text{SSIM}} + \ell_{\text{DC}}$

- **Categorical Cross Entropy (CCE)-pixel-level:**

$$\ell_{\text{CCE}} = - \sum_{i=1}^C \sum_{a=1}^H \sum_{b=1}^W y_{(a,b)}^i \ln y_{*(a,b)}^i,$$

— — —multi-class classification and segmentation.

- **Structural Similarity (SSIM)-patch-level:**

$$\ell_{\text{SSIM}} = 1 - ((2\mu_S\mu_G + \varepsilon_1)(2\sigma_{SG} + \varepsilon_2)) / ((\mu_S^2 + \mu_G^2 + \varepsilon_1)(\sigma_S^2 + \sigma_G^2 + \varepsilon_2)),$$

— — —assessing image quality and is used to capture the structural information.

- **Dice Coefficient (DC)-map-level:**

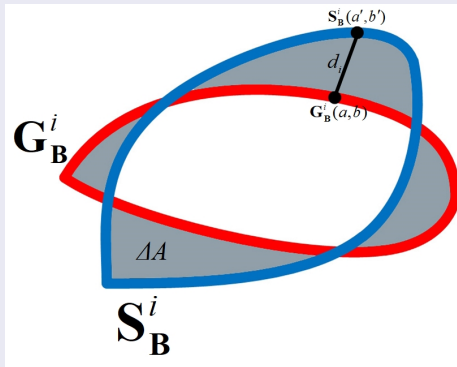
$$dice_i = (\epsilon + 2 \sum_{n=1}^{N_i} y_n^i y_{*n}^i) / (\epsilon + \sum_{n=1}^{N_i} (y_n^i + y_{*n}^i));$$

$$\ell_{\text{DC}} = 1 - \sum_{i=1}^C dice_i / (N_i + \epsilon)$$

— — —measuring the similarity between two sets.

The region loss guides the network to learn the transformation **in a three-level hierarchy**.

$$\text{Boundary Loss: } \ell_B = \sum_i^C \int_{G_B^i} \|S_B^i(a', b') - G_B^i(a, b)\|^2 d(a, b)$$



The boundary loss function is to optimize the segmentation result.

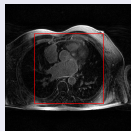
# Experiments and Results

## Dataset description

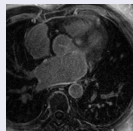
- 100 annotated 3D MRIs
  - — —80 cases for training and 20 cases for validating.
- Pixel spacing:  $0.625 \times 0.625 \times 0.625$  mm/pixel.
- Image sizes:  $88 \times 576 \times 576$  and  $88 \times 640 \times 640$  pixels.

## Pre-processing

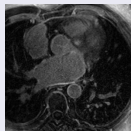
- Cropping each slice to  $346 \times 346$  pixels.
- Gaussian normalization.
- Stacking 3 successive 2D frames.



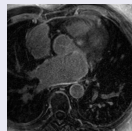
Initial image



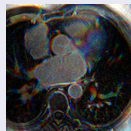
$n - 1$



$n$



$n + 1$



"3D-Like"

## Post-processing

- Keeping the greatest connected component of the segmentation.
- Padding with zeros to get back a initial width and height of a slice.

## Implementation and Experimental Setup

- Keras/TensorFlow using a Nvidia Quadro P6000 GPU.
- Hybrid loss function
  - softmax to get a probability distribution over classes.
- Adam optimizer
  - batchsize=3,  $\beta_1=0.9$ ,  $\beta_2=0.999$ , epsilon=0.001, lr = 0.01.
- Epoch=30.
- Evaluation method:
  - Dice coefficient;
  - 95% Hausdorff distance (95HD);
  - Average Hausdorff distance (AHD).

# Results on training datasets

Comparison of our method and other state-of-the-art architectures using a 5 fold cross-validation.

Method	Att. Module	Hyb. Loss	DC/%	95HD/mm	AHD/mm
U-Net [3]			88.556( $\pm 2.586$ )	4.447( $\pm 0.996$ )	0.212( $\pm 0.077$ )
		✓	89.613( $\pm 2.257$ )	4.169( $\pm 0.960$ )	0.210( $\pm 0.118$ )
DANet [2]			84.229( $\pm 3.774$ )	6.145( $\pm 2.341$ )	0.514( $\pm 0.477$ )
		✓	87.584( $\pm 2.765$ )	4.903( $\pm 1.448$ )	0.280( $\pm 0.179$ )
Deeplabv3+ [1]			85.444( $\pm 3.079$ )	5.872( $\pm 2.345$ )	0.504( $\pm 0.614$ )
		✓	87.556( $\pm 1.155$ )	5.210( $\pm 1.087$ )	0.273( $\pm 0.074$ )
Our Method			90.774( $\pm 1.568$ )	3.312( $\pm 1.277$ )	0.158( $\pm 0.092$ )
	✓		91.326( $\pm 1.174$ )	3.097( $\pm 0.810$ )	0.143( $\pm 0.055$ )
	✓	✓	<b>91.792(<math>\pm 1.065</math>)</b>	<b>2.868(<math>\pm 0.667</math>)</b>	<b>0.130(<math>\pm 0.042</math>)</b>



Chen, L.C., et al.: Encoder-decoder with atrous separable convolution for semantic image segmentation. In: Proc. of ECCV. pp. 801–818 (2018)



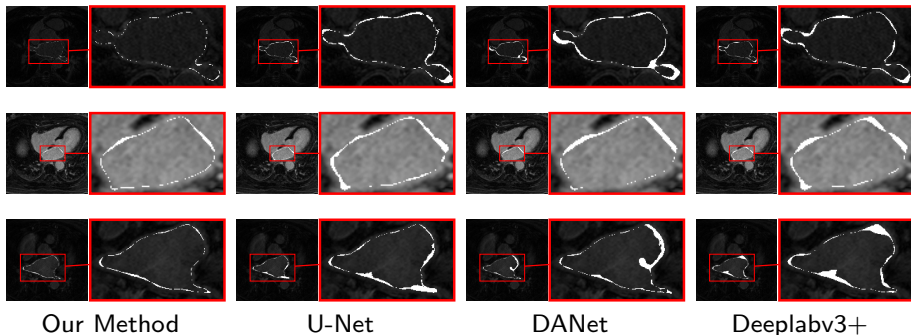
Fu, J., et al.: Dual attention network for scene segmentation. In: Proc. of CVPR. pp. 3146–3154 (2019)



Ronneberger, O., Fischer, P., Brox, T.: U-Net: Convolutional networks for biomedical image segmentation. In: Proc. of MICCAI. LNCS, vol. 9351, pp. 234–241. Springer (2015)



Comparison of the proposed method and other state-of-the-art architectures. The white pixels are the differences between the prediction and the GT.



- **Treat boundaries and regions fairly**
- **Hybrid loss**
  - considering regions and boundaries of objects equally by combining region loss with boundary loss.
- **Attention module**
  - preventing the interferences between the surrounding similar tissues and to capture large-scale and thinner structures
- **Temporal-like method**
  - taking advantage of the temporal information by stacking 3 successive 2D frames.

# Thank you!

