

# Temporal Image Registration using deep learning for 3D Fetal Echocardiography

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### **Outline**

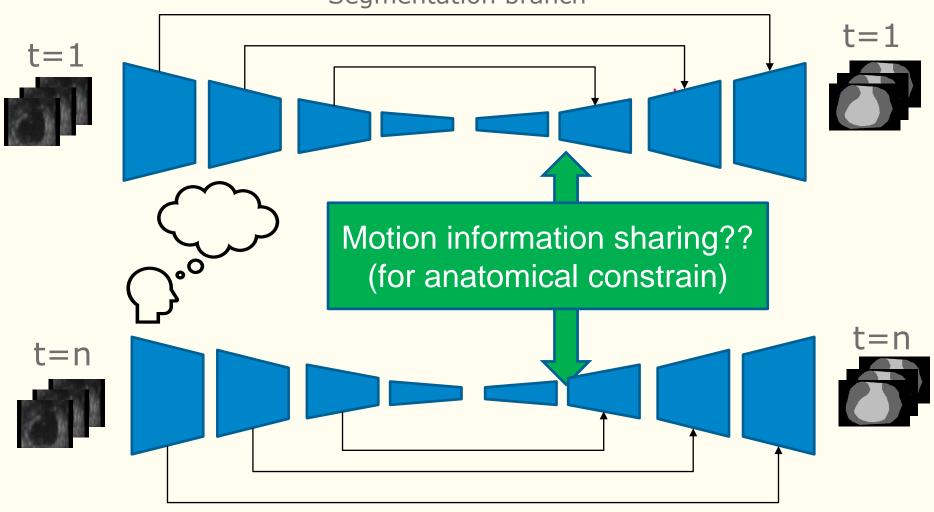
- Introduction
- State of the Art
- Dataset description
- Proposed Registration Framework
- Results
- Conclusion & Future Remarks
- References

## Introduction

- The fetal heart can experience congenital heart malformation and functional abnormalities.
- Ultrasound imaging plays a vital role in assessing the heart of the developing fetus due to its non-invasive nature.
- Heart chambers, valves, blood flow patterns, etc. can be used as good identifiers to detect and evaluate several cardiac diseases.
- Mowever, the detection of heart problems in fetus via mass screening is only around 50%
- The clinical use of echo is still stuck with 2D
- Most of the works are based on adult hearts

# **Temporal Image Registration**

Segmentation branch



Segmentation branch

The estimation of the **deformation field** by registration between two time points can help share the information between two segmentation branches [1][2][3].

- [1] Gupta, Soumya, et al. "Multi-class motion-based semantic segmentation for ureteroscopy and laser lithotripsy." *Computerized Medical Imaging and Graphics* 101 (2022): 102112.
- [2] Gupta, Soumya, et al. "Mi-unet: Improved segmentation in ureteroscopy." 2020 IEEE 17th International Symposium on Biomedical Imaging (ISBI). IEEE, 2020.
- [3] Xue, Wufeng, et al. "Improved Segmentation of Echocardiography With Orientation-Congruency of Optical Flow and Motion-Enhanced Segmentation." IEEE Journal of Biomedical and Health Informatics 26.12 (2022): 6105-6115.

# Imperial College State Of the Art (Dataset)

- Public Dataset for 3D echocardiography is very rare.
- 2 very well known public datasets are:
  - o **CAMUS** [5]:
    - the largest publicly-available and fully-annotated
    - > Limitations: data is **2D** and from **adult hearts**
  - o **CETUS** [14]: 45 **3D** echocardiographic sequences
    - Adult hearts
  - Nurmaini et. Al. [13] worked on a private dataset of 1149
     2D images for fetal hearts.

# Imperial College London State Of the Art (Methods)

\*Dice for Intra observer variability **0.930** 

Work	Description	Result ( Mean Dice Score)
Smistad et al. [15] 2017	<ul> <li>used <b>U-Net CNN</b> to segment the left ventricle in 2D ultrasound images.</li> </ul>	<b>0.87</b> on LV (manual annotated test set)
Oktay et al. [16] 2019	<ul> <li>used anatomically constrained neural network (ACNN) segment the 3D LV structure .</li> <li>Uses segmentation aware maps to add anatomically constraints</li> </ul>	<b>0.912</b> (ED) and <b>0.873</b> (ES) on <b>CETUS</b> dataset
Wei et al. [18] 2020	<ul> <li>proposed appearance and shape level co-learning</li> <li>mutual benefits of the segmentation and tracking</li> </ul>	<b>0.929</b> for LV and Myo on <b>CAMUS</b> dataset

# Imperial College London State Of the Art (Methods)

\*Dice for Intra observer variability **0.930** 

Work	Description	Result ( Mean Dice Score)
Zhang et al. [18] 2022	<ul> <li>dual-branch TransV-Net (DBTV)</li> <li>V-shaped encoder-decoder branches</li> <li>extract the additional edge features</li> </ul>	<ul><li>0.913 and 0.880 for left and right ventricle segments on CAMUS Dataset</li></ul>
Sfakianakis et al. [17] 2023	<ul> <li>used ensemble of CNNs based on the U-net architecture</li> <li>Incorporated geometrically constrained data augmentation method based</li> </ul>	<b>0.929</b> on LV and endocardium on <b>CAMUS</b> Dataset
Ling et al. [19] 2023	<ul> <li>proposed four models based on nnUNet (53M parameters)</li> <li>data augmentation in both training and inference, combined with a well-matched optimization scheme</li> </ul>	<ul><li>0.935 crossing Intra</li><li>observer variability dice</li><li>score for LV and myo on</li><li>CAMUS dataset</li></ul>

# Fetal Data Annotation (3D)



📋 t1.avi	24/7/2019 10:14 AM	AVI File	52,829 KB 00:00:02
🖹 t2.avi	24/7/2019 10:16 AM	AVI File	52,829 KB 00:00:02
🖹 t3.avi	24/7/2019 10:19 AM	AVI File	52,829 KB 00:00:02
🖆 t4.avi	24/7/2019 10:26 AM	AVI File	52,829 KB 00:00:02
🖹 t5.avi	24/7/2019 10:26 AM	AVI File	52,829 KB 00:00:02
🖹 t6.avi	24/7/2019 10:27 AM	AVI File	52,829 KB 00:00:02
🖹 t7.avi	24/7/2019 10:28 AM	AVI File	52,829 KB 00:00:02
📋 t8.avi	24/7/2019 10:29 AM	AVI File	52,829 KB 00:00:02
🖆 t9.avi	24/7/2019 10:33 AM	AVI File	52,829 KB 00:00:02
🖹 t10.avi	24/7/2019 10:34 AM	AVI File	52,829 KB 00:00:02
🖆 t11.avi	24/7/2019 10:34 AM	AVI File	52,829 KB 00:00:02
📋 t12.avi	24/7/2019 10:36 AM	AVI File	52,829 KB 00:00:02
🖹 t13.avi	24/7/2019 10:37 AM	AVI File	52,829 KB 00:00:02
📋 t14.avi	24/7/2019 10:37 AM	AVI File	52,829 KB 00:00:02
📋 t15.avi	24/7/2019 10:38 AM	AVI File	52,829 KB 00:00:02
🖹 t16.avi	24/7/2019 10:39 AM	AVI File	52,829 KB 00:00:02
📋 t17.avi	24/7/2019 10:39 AM	AVI File	52,829 KB 00:00:02
🖹 t18.avi	24/7/2019 10:40 AM	AVI File	52,829 KB 00:00:02
🖹 t19.avi	24/7/2019 10:41 AM	AVI File	52,829 KB 00:00:02
🖹 t20.avi	24/7/2019 10:42 AM	AVI File	52,829 KB 00:00:02
🗌 🖹 t21.avi	24/7/2019 10:42 AM	AVI File	52,829 KB 00:00:02
🖹 t22.avi	24/7/2019 10:43 AM	AVI File	52,829 KB 00:00:02
🖹 t23.avi	24/7/2019 10:44 AM	AVI File	52,829 KB 00:00:02

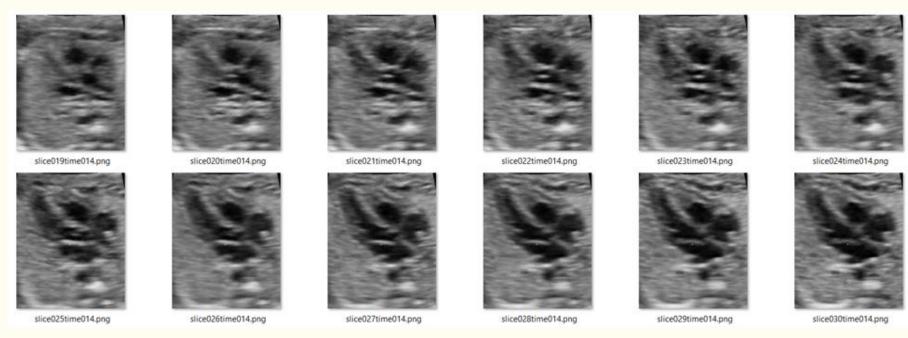
Calculate cine at each time point

- 1. With same cine length (adjust start and end slice until cannot see ventricle)
- 2. Step size: 0.5mm
- 3. Press Start



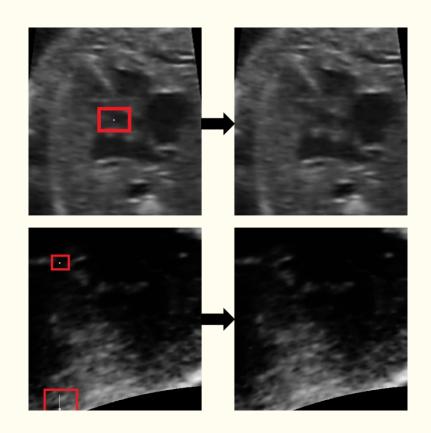
#### **Generate Slices: AVI to PNG**

```
Editor - E:\Programming Code\MatlabCode\extract_time.m
   extract_time.m × generate_fourier_stls.m × View_across_time.m × +
13
       %%%%%%%%% interest
       %%%%%%%% - right click within the box and click "crop image"
14
15
16 -
       bar_dist=10; % gauge bar distance of the two points, chosen to be 10mm
17 -
       TUI dist=0.5; % please change the TUI dist
       max time=36;
        largest time=34;
19 -
       video_dir='E:\healthy_fetus_heart_case\01082019';
21
22 -
       Obj=VideoReader([video_dir '\video\' sprintf('t%d.avi',largest_time)]);
       vid=read(Obj);
```



## **Preprocessing**

- The ultrasound intensity images contains some artifacts like constant white boxes or arrows
- linear interpolation method was used to remove them
- defected area was interpolated using the intensity values from the interpolation line



# **Image Registration**

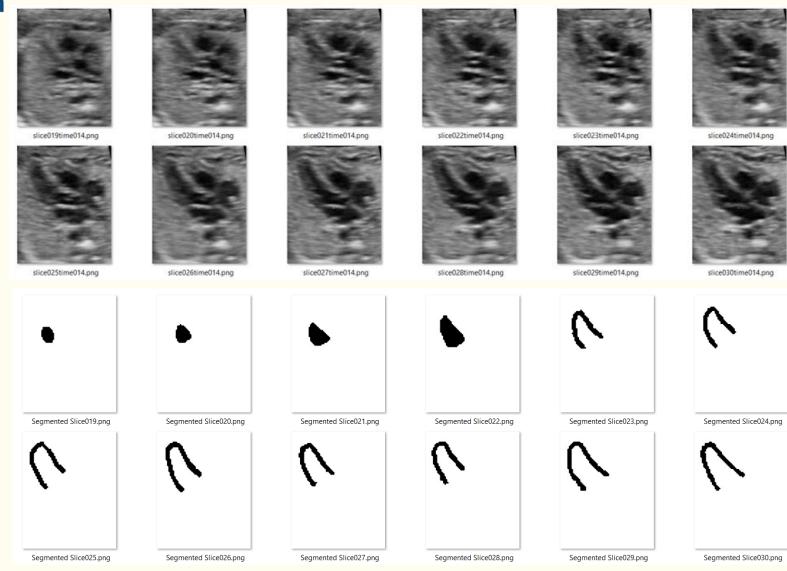
```
(base) E:\healthy_fetus_heart_
medImgProc version 1.0.0
no display found. Using non-ir
Registering t 2 wrt t 1
Registering t 2 wrt t 0
Registering t 3 wrt t 2
Registering t 3 wrt t 0
Registering t 4 wrt t 3
Registering t 4 wrt t 0
Registering t 5 wrt t 4
Registering t 5 wrt t 0
Registering t 6 wrt t 5
Registering t 6 wrt t 0
Registering t 7
                wrt t 6
Registering t 7
                wrt t 0
Registering t 8
                wrt t 7
Registering t 8 wrt t 0
Registering t 9
                wrt t 8
```

- Register image with respect to t1 and the previous time point
- Libraries used:
  - SimpleElastix
  - Cardiac motion estimation
    library by Wiputra et al.
     (2020)[1]
    [https://github.com/WeiXuanChan/motionSegmentation]
- uses the Fourier b-splines spatiotemporal motion model to fit the deformation fields.

# **Image Segmentation (lazysnap)**

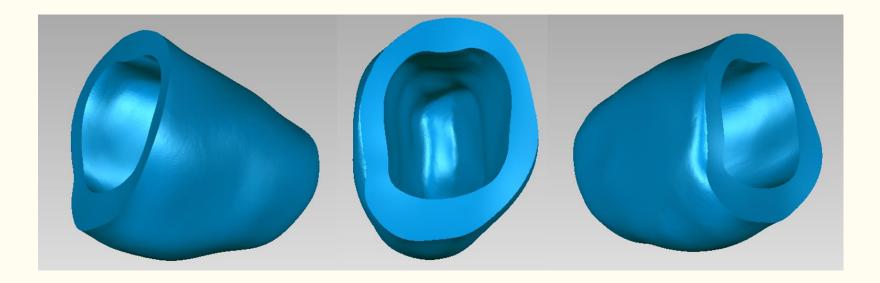
```
Give the path to the folder where folders containing images are stored:
E:\healthy_fetus_heart_case\01082019
Give the number of Slices:
Give the number of Times:
27
Give start time:
27
Give start slice:
Give image skip value (default 0, no skip):
0
E:\healthy_fetus_heart_case\01082019/time027/segmented/Segmented Slice006.png
Press 'f'then hold and drag left click to select the foreground region
Press 'b'then hold and drag left click to select the background region
Press 'r' to reset the image segmentation procedure
Press esc to quit and safe the images
```

- ☐ **Two** time points to segment
  - 1. End-systolic
  - 2. Fnd-diastolic

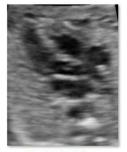


#### **3D Reconstruction**

- vmtk(Vascular Modeling Toolkit) was used to generate the 3D masks
- Generated 3D masks have artifacts like holes, spikes, edges etc which was corrected and smoothened using Geomagic by an expert.



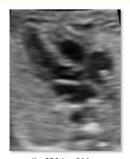
# **Image-Mask Pair**



slice025time014.png



Segmented Slice025.png



slice026time014.png



Segmented Slice026.png



slice027time014.png



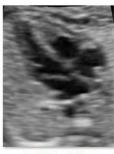
Segmented Slice027.png



slice028time014.png



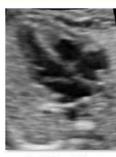
Segmented Slice028.png



slice029time014.png



Segmented Slice029.png



slice030time014.png



Segmented Slice030.png

## **Fetal 3D Dataset**

H: Healthy

D: Diseased





Train: 10 Cases Val: 4 Cases (2-H, 8-D)



381 3D Volumes 123 3D Volumes Pairs of fixed and moving

(2-H, 2-D)

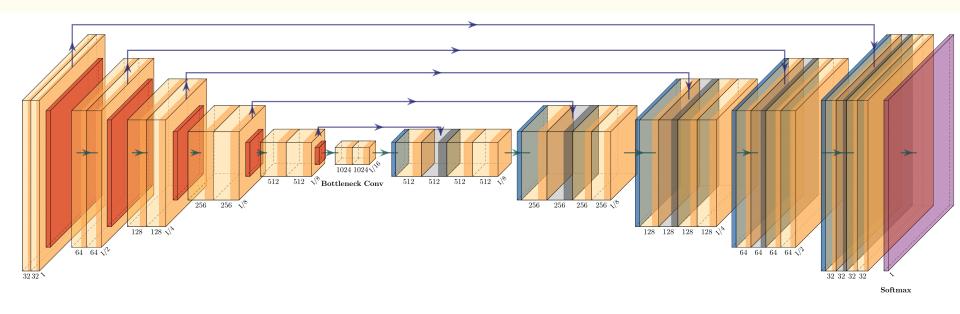
Pairs of fixed and moving

## **CAMUS 2D Dataset**

- We have also applied our experimental methods on a well-known adult 2D echocardiography dataset.[5]
- We wanted to compare the performances in 2D vs 3D
- Also in adult vs fetal echocardiography
- CAMUS dataset comprises :
  - i) a training set of **450** patients along with the corresponding manual references;
  - ii) a testing set composed of **50** new patients.

# **Proposed registration Framework**

# (1) Proposed Residual Segmentation branch



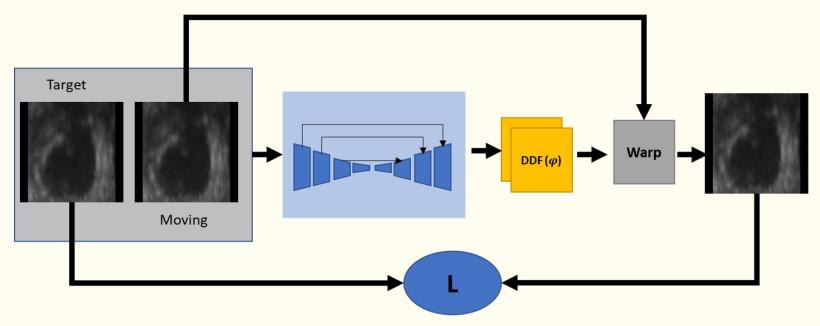
- Based on the traditional **UNET** architecture with skip connections. (Input Image: 256\*256\*32)
- The conventional path cannot degrade the features' quality as a non-zero regularizing path will skip over them. On the other hand, the direct skipping of the non-zero regularizing path cannot hamper the performance as it has been added to the conventional path's learned features.

(2) Multi-class Anatomically Constrained and multi-scale registration

#### The proposed registration method has the following integral parts:

- 1. Local and global anatomical constraints using variational autoencoders
- Adversarial learning as like zero-sum game theory (one agent's gain is another agent's loss), where the discriminator is used to classify moved and fixed images
- 3. Multi-scale (multi-resolution) training, where trained parameters in lower scale are used to initialize the higher scale training (Future works for 3D)

#### Vanilla-DLIR



Typical unsupervised Registration without considering anatomy.

$$\mathcal{L}_{us}(f, m, d) = \mathcal{L}_{sim}(f, m \circ d) + \lambda \mathcal{L}_{smooth}(d)$$

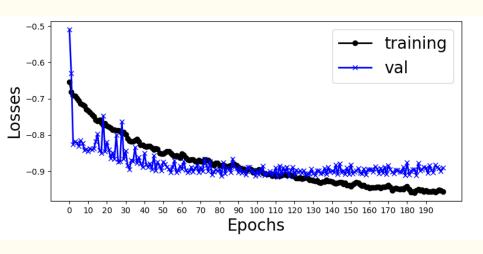
$$\text{MI}(f_x; m_y) = \sum_{x \in X} \sum_{y \in Y} P(f_x, m_y) \log_2 \left(\frac{P(f_x, m_y)}{P(f_x)P(m_y)}\right)$$

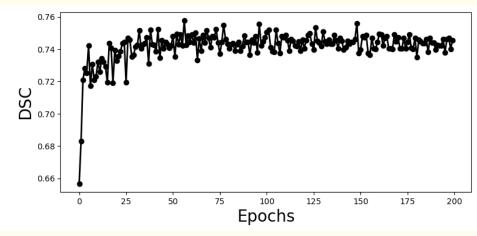
$$\mathcal{L}_{smooth}(d) = \sum_{\mathbf{d} \in D} \| \nabla \mathbf{u}(\mathbf{d}) \|^2$$

 The unsupervised loss can be derived using MI loss and binding energy regularization loss

#### Vanilla-DLIR

#### Results on Fetal Dataset (3D) – Longitudinal Registration



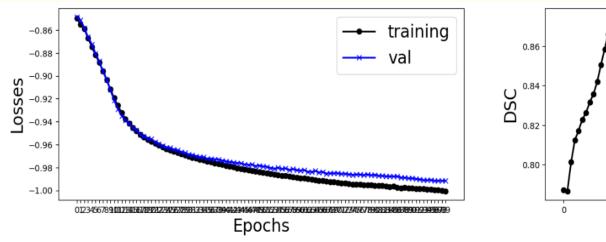


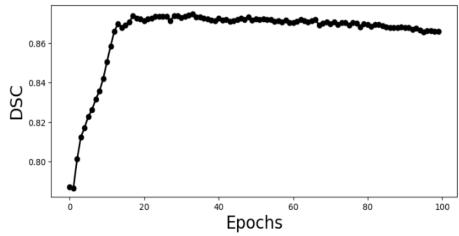
MSEMetric W/O registration = 0.00377 MSEMetric W/ registration = 0.00296

BG LV Myo Mean dice W/O registration =  $[0.99093 \ 0.78917 \ 0.72605]$ Mean dice W/ registration =  $[0.98699 \ 0.70087 \ 0.58543]$ 

#### **Vanilla-DLIR**

#### Results on CAMUS Dataset (2D) - Longitudinal Registration

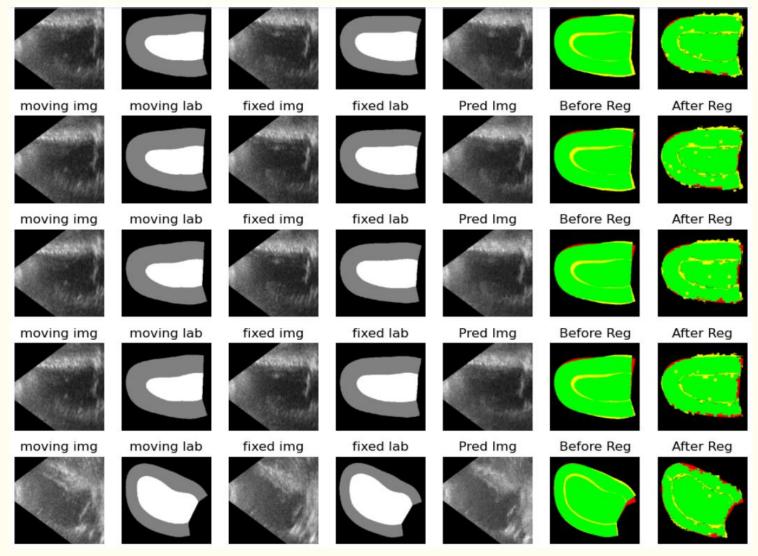




MSEMetric W/O registration = 0.00972 MSEMetric W/ registration = 0.0042

BG LV Myo Mean dice W/O registration = [0.96678 0.76046 0.69391] Mean dice W/ registration = [0.97352 0.87523 0.74977]

#### Vanilla-DLIR



#### Vanilla-DLIR

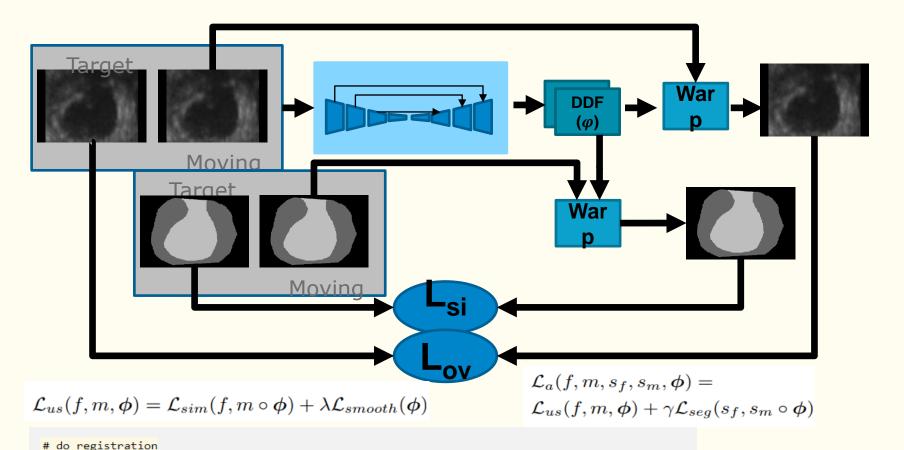
#### **Observation:**

 In the unsupervised registration without considering the anatomy, the similarity between two intensity images increases, but the similarity between fixed and moved masks does not improve satisfactorily or fail in some cases.

# Thanks to those two following papers that solved this limitation.

[1] Balakrishnan, Guha, et al. "VoxelMorph: a learning framework for deformable medical image registration." IEEE transactions on medical imaging 38.8 (2019): 1788-1800.

[2] Hu, Yipeng, et al. "Label-driven weakly-supervised learning for multimodal deformable image registration." 2018 IEEE 15th International Symposium on Biomedical Imaging (ISBI 2018). IEEE, 2018.



```
displacement_field = reg_net(torch.cat((img1,img2), dim=1))

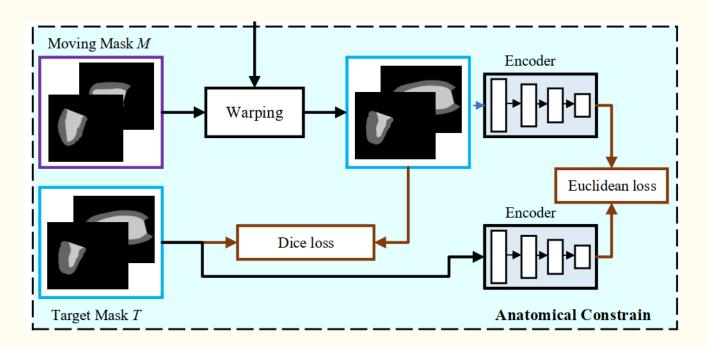
# do segmentation
seg1 = seg_net(img1).softmax(dim=1)
seg2 = seg_net(img2).softmax(dim=1)

# warp segmentation using the same warp block defined above
seg2_warped = warp(seg2, displacement_field)

# compute multiclass dice loss
dice_loss = monai.losses.DiceLoss()
anatomy_loss = dice_loss(seg2_warped, seg1)
```

•  $\mathcal{L}_{seg}$  is measured using the **Dice-Loss**.

# Imperial College Rationale for Latent Space consideration London

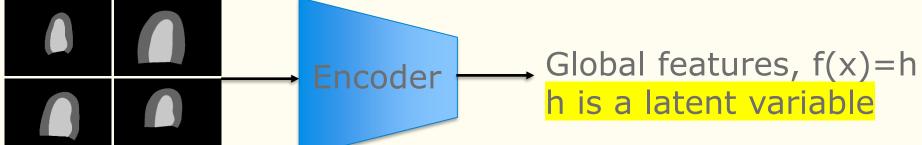


#### To improve the network's anatomical context:

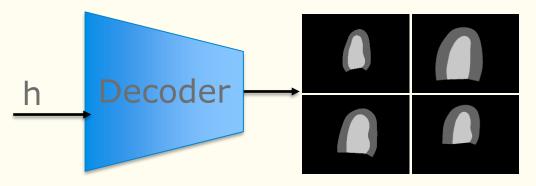
Intensity-based similarity loss (fixed and moving intensity images) with a segmentation-aware loss (fixed and moved labels) are combined.

This new loss measures the alignment between a target anatomical mask and a warped moving mask.

# **Learning Global Anatomical Features**



Input Shapes (x)



#### Loss

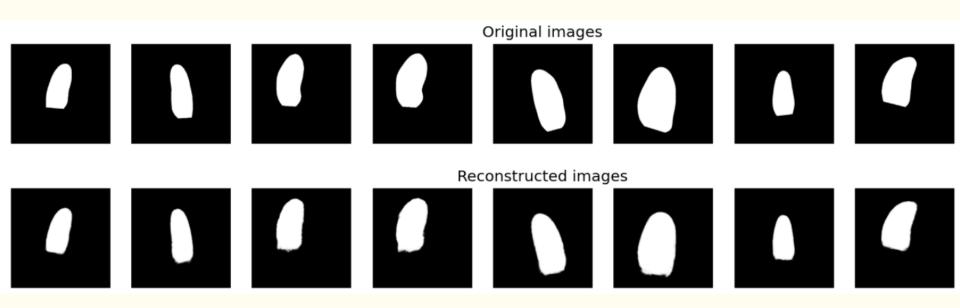
- Cross-entropy loss (x, y)
- 2. KL loss
- 3. [NEW] Similarity loss (x, y) like DSC, **SSIM** etc

Reconstructed Shapes (y)

The regularization loss named **Kullback-Leibler (KL)** divergence forces the distributions returned by the encoder to be close to a standard normal distribution.

## **Goal from Global attributes**

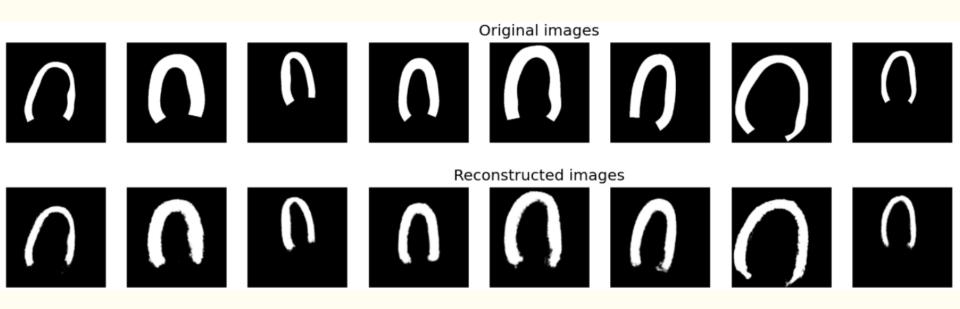
Orientation, Shape, and Size



Best Val Dice = **0.9789** 

## **Goal from Global attributes**

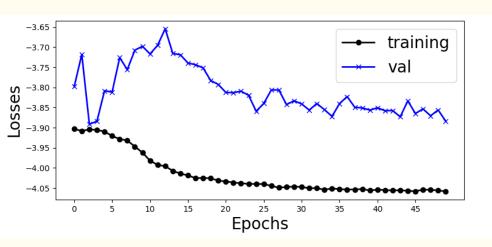
Orientation, Shape, and Size

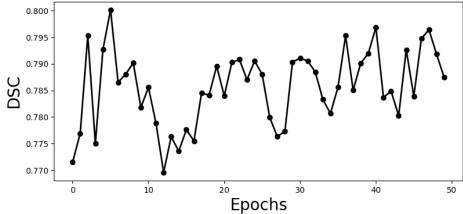


Best Val Dice = **0.9407** 

#### **AC-DLIR**

#### Results on Fetal Dataset (3D) – Longitudinal Registration



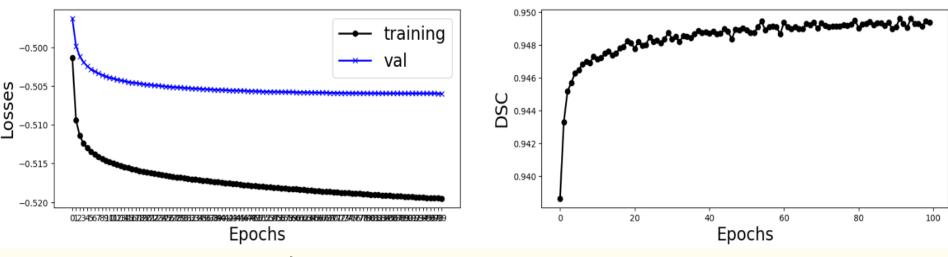


MSEMetric W/O registration = 0.00377 MSEMetric W/ registration = 0.00251

BG LV Myo Mean dice W/O registration =  $[0.99093 \ 0.78917 \ 0.72605]$ Mean dice W/ registration =  $[0.98959 \ 0.73347 \ 0.64435]$ 

#### **AC-DLIR**

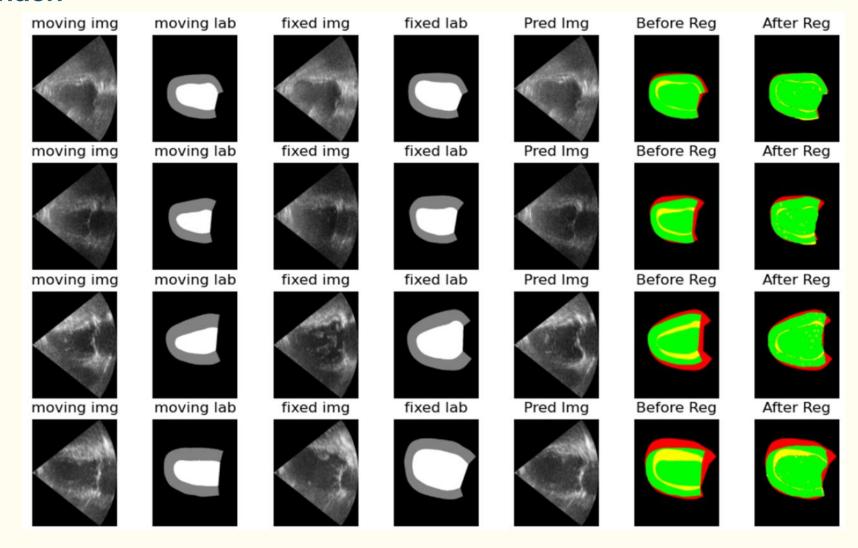
#### Results on CAMUS Dataset (2D) - Longitudinal Registration



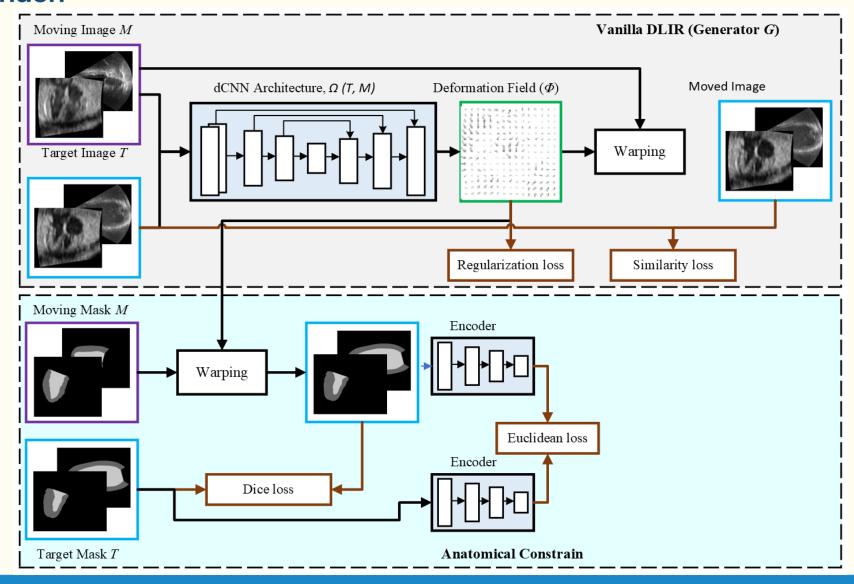
MSEMetric W/O registration = 0.00972 MSEMetric W/ registration = 0.00598

BG LV Myo Mean dice W/O registration =  $[0.96678 \ 0.76046 \ 0.69391]$ Mean dice W/ registration =  $[0.97972 \ 0.91935 \ 0.81437]$ 

#### **AC-DLIR**



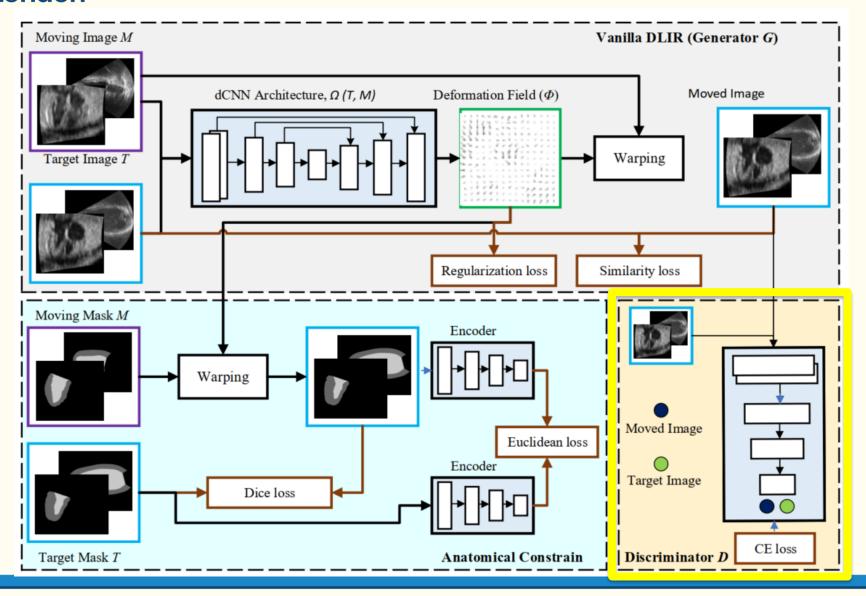
# Imperial College Anatomically constrained DLIR (AC-DLIR) London



# Imperial College Adversarial AC-DLIR (AdvAC-DLIR) London

- Still, there is room for performance improvement.
- Hence, we proposed Adversarial Anatomically constrained (AdvAC) DLIR framework.
- The part of AC-DLIR for generating the deformable images as a generator for the adversarial network.
- A discriminator was also trained which was able to classify the fixed and moved images
- While training, the generator would try to create as much as
   plausible images as the fixed image whereas the discriminator
   would try to discriminate them.

# Imperial College Adversarial AC-DLIR (AdvAC-DLIR) London



# **Proposed Loss Functions**

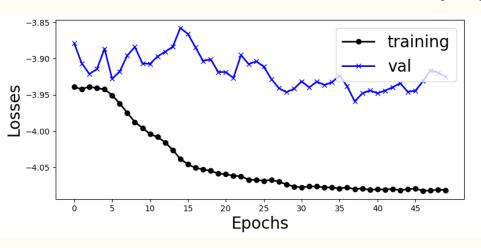
$$\mathcal{L}_{advac}(f, m, r_f, r_m, d, s_m) = \mathcal{L}_{us}(f, m, d) + \beta \mathcal{L}_{dice}(r_f, r_m \circ d) + \gamma \mathcal{L}_{L2}(r_f, r_m) + \mathbf{\phi} \mathcal{L}_{\mathbf{g}}(\mathbf{m}, \mathbf{s}_{\mathbf{m}})$$

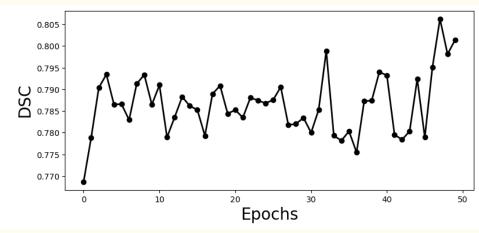
# Here,

- $\mathcal{L}_{us}$  is the unsupervised loss function coming from Vanilla-DLIR.
- $\mathcal{L}_{dice}$  is the segmentation aware loss
- $\mathcal{L}_{L2}$  is the loss from latent space consideration
- $\mathcal{L}_{g}$  is the generator's binary cross entropy loss
- and  $\beta$ ,  $\gamma$  and  $\phi$  are regularization parameters

# **AdVAC-DLIR**

# Results on Fetal Dataset (3D) – Longitudinal Registration



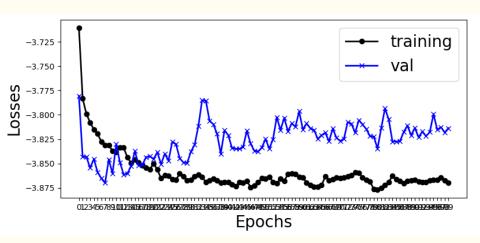


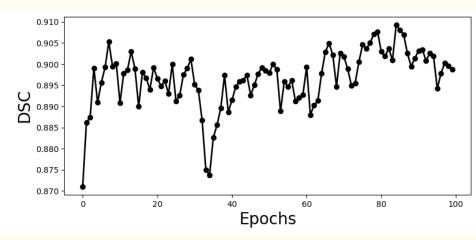
MSEMetric W/O registration = 0.00377 MSEMetric W/ registration = 0.00258

BG LV Myo Mean dice W/O registration =  $[0.99093 \ 0.78917 \ 0.72605]$ Mean dice W/ registration =  $[0.99089 \ 0.79884 \ 0.73482]$ 

# **AdVAC-DLIR**

# Results on CAMUS Dataset (2D) - Longitudinal Registration

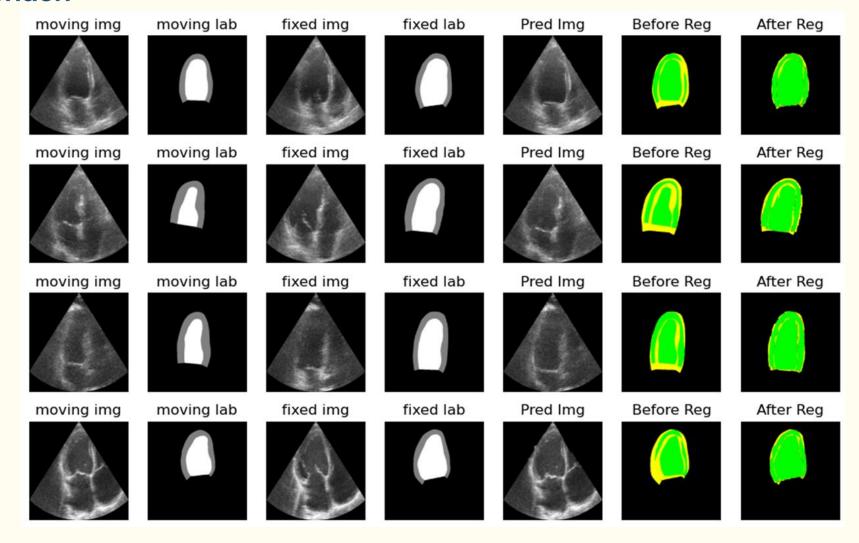




MSEMetric W/O registration = 0.00972 MSEMetric W/ registration = 0.00589

BG LV Myo Mean dice W/O registration =  $[0.96678 \ 0.76046 \ 0.69391]$  Mean dice W/ registration =  $[0.98742 \ 0.93573 \ 0.82751]$ 

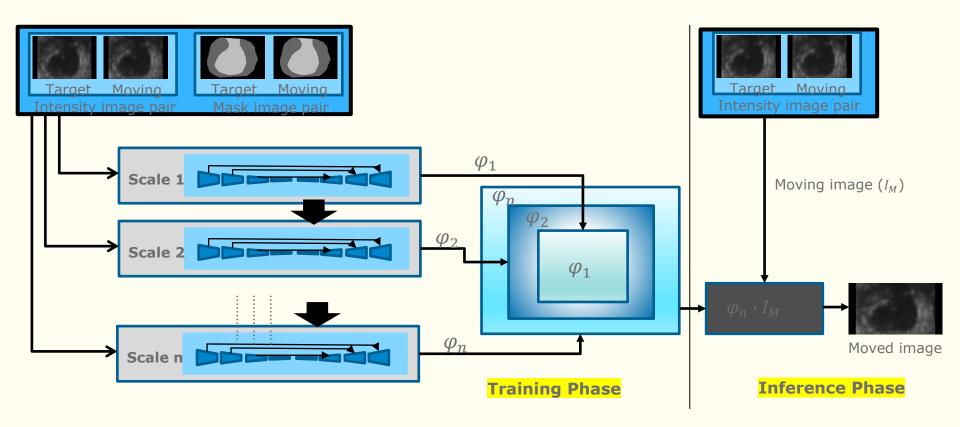
# **AdVAC-DLIR**



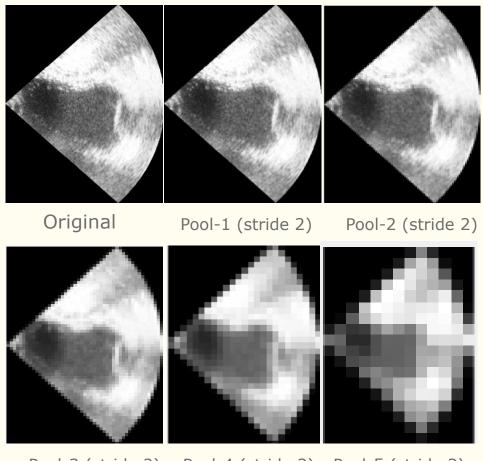
# Results (Fetal 3D)

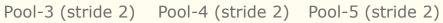
Model	MSE	Dice Score			Mean Dice
		Background	LV	Myo	±std
Without Registration	0.00377	0.99093	0.78917	0.72605	$0.83539 \pm 0.12798$
Vanilla- DLIR	0.00296	0.98699	0.70087	0.58543	0.75776± 0.04036
AC-DLIR	0.00251	0.98959	0.73347	0.64435	0.80013± 0.05401
Adv-DLIR	0.00339	0.99031	0.73836	0.67389	0.80989± 0.05142
AdvAC- DLIR	0.00258	0.99089	0.79884	0.73482	0.84668± 0.04586

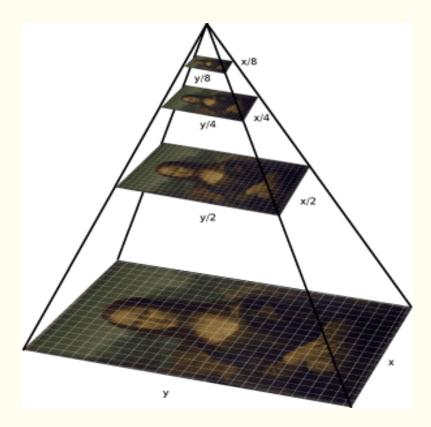
# **Future Plan : MACMR framework**



Multi-scale (multi-resolution) training, where trained parameters
 in lower scale are used to initialize the higher scale training







#### Results on CAMUS dataset

```
Scale 1 (112 \times 144)
```

```
before_MSEMetric = 0.010041479021310806
after_MSEMetric = 0.006580408196896315
before_compute_meandice = [0.9668386  0.6937522  0.75969636]
after_compute_meandice = [0.9774292  0.7890314  0.9059601]
```

# Scale 2 $(224 \times 288)$

```
before_MSEMetric = 0.009544618427753448

after_MSEMetric = 0.005924441386014223

before_compute_meandice = [0.9667753  0.69394  0.76024437]

after_compute_meandice = [0.97944987  0.8144262  0.9186317 ]
```

```
Scale 3 (448 \times 576)
```

```
before_MSEMetric = 0.009694118052721024

after_MSEMetric = 0.0059749712236225605

before_compute_meandice = [0.96685976 0.69459504 0.76026726]

after_compute_meandice = [0.9799058 0.81560487 0.9229658 ]
```

# Results (CAMUS 2D)

Model	MSE	Dice Score			Mean Dice
		Background	Myo	LV	±std
Without Registration	0.00972	0.96678	0.69391	0.76046	0.80235± 0.05491
Vanilla- DLIR	0.0042	0.97352	0.74977	0.87523	0.88487± 0.03261
AC-DLIR	0.00598	0.97972	0.81437	0.91935	0.90303± 0.03447
Adv-DLIR	0.00533	0.97429	0.79278	0.86842	0.85733± 0.04129
AdvAC- DLIR	0.00589	0.98742	0.82751	0.93573	0.91689± 0.02596
MACMR	0.00489	0.98779	0.84871	0.95423	0.94245± 0.02474

# Conclusion

- The clinical use of echo is still stuck with 2D, so a more robust and precise automatic registration and segmentation framework is really needed
- Proposed AdvAC model outperforms all the previous experiments and thus proved to be the best model working in both the 2D and 3D dataset.
- Still, there is room for performance improvement.
- More annotated data for 3D will be needed for further performance improvement.

# **Future Plans**

- Annotate more cases for 3D fetal data
- Apply the multi-resolution (MACMR) framework for 3D data
- Proceed with the 3D segmentation of heart chamber and myocardium
- Measure various identifiers as heart shapes, ejection fraction, volume etc. to perform disease detection and classification for fetal heart abnormalities

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# Thank you

Do you have any questions?