

Spatiotemporal Neural Diffeomorphic Flow for Cardiac Atlas and Motion Recovery

Yiyang Xu¹

YIYANG.1.XU@KCL.AC.UK

¹ *School of Biomedical Engineering and Imaging Sciences, King’s College London, UK*

Esther Puyol-Antón²

EPUYOLANTON@HEARTFLOW.COM

Matthew Sinclair²

MSINCLAIR@HEARTFLOW.COM

² *Heartflow Inc., Mountain View, USA*

Amedeo Chiribiri¹

AMEDEO.CHIRIBIRI@KCL.AC.UK

Steven A Niederer³

STEVEN.NIEDERER@IMPERIAL.AC.UK

³ *National Heart and Lung Institute, Imperial College London, UK*

Alistair A Young¹

ALISTAIR.YOUNG@KCL.AC.UK

Editors: Under Review for MIDL 2026

Abstract

Spatiotemporal cardiac atlases built from cardiovascular magnetic resonance (CMR) data enable population-level analysis of cardiac anatomy and motion, yet existing methods lack topological guarantees and dense inter-subject correspondences. We extend Neural Diffeomorphic Flow (NDF) to the spatiotemporal domain, constructing a population-based left ventricular atlas with dense, topology-preserving correspondences across individuals and time frames. Evaluated on UK Biobank CMR data, our method achieves 1.4 mm chamfer distance for motion recovery of the full cardiac cycle from a single end-diastolic frame.

Keywords: Cardiovascular Magnetic Resonance, Spatiotemporal Cardiac Atlases, Motion Recovery

1. Introduction

Cardiovascular diseases remain the leading cause of mortality worldwide (Mendis et al., 2011), driving demand for computational approaches that quantify cardiac structure and dynamics non-invasively. Cardiovascular magnetic resonance (CMR) is the gold standard for capturing full-cycle cine sequences at high spatiotemporal resolution without ionising radiation (Petersen et al., 2016). Computed tomography (CT) offers finer spatial detail but is limited to a single cardiac phase due to dose constraints. Spatiotemporal cardiac atlases establish shared reference spaces for cross-subject comparison and disease characterisation (Young and Frangi, 2009), yet classical approaches (Bai et al., 2015) require costly registration and fixed discretisations, while methods such as CHeart (Qiao et al., 2024) and CardiacFlow (Ma et al., 2025) learn implicit atlases via latent spaces but lack topological guarantees and the ability to map between atlas and individual. Neural Diffeomorphic Flow (NDF) (Sun et al., 2022) addresses these limitations by parameterising shape-conditioned deformations as neural ODE solutions, ensuring invertibility and topology preservation by construction. We extend NDF to the spatiotemporal domain, learning diffeomorphic flows that capture shape and motion variations across subjects and time, yielding a left ventricular atlas with dense, topology-preserving correspondences to individual cases. We validate

on UK Biobank CMR data (Sudlow et al., 2015), demonstrating effective motion recovery from sparse frames while preserving anatomical plausibility. Our contributions are: (1) extending NDF to 4D for cardiac motion modelling; (2) constructing an explicit spatiotemporal left ventricular atlas with topology-preserving correspondences; and (3) validating motion recovery on UK Biobank data with ground-truth evaluation.

2. Methods

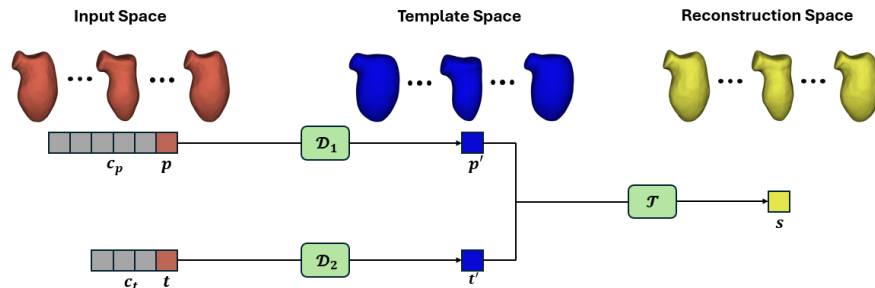


Figure 1: Overview of spatiotemporal NDF. A spatial warper \mathcal{D}_1 and temporal warper \mathcal{D}_2 map input spatial and temporal coordinates (\mathbf{p}, t) with subject-specific latent codes $(\mathbf{c}_p, \mathbf{c}_t)$ to template coordinates (\mathbf{p}', t') , which are passed to a single-shape DeepSDF (Park et al., 2019) \mathcal{T} to predict signed distance s .

Rather than operating on 3D coordinates alone, we extend NDF to the spatiotemporal domain with two independent deformation modules for the spatial and temporal dimensions. Given a CMR sequence X_i , the network accepts (\mathbf{p}, t) and a subject-specific latent code \mathbf{c}_i as input, where $t \in [0, 1]$ denotes the normalised cardiac cycle phase and \mathbf{p} denotes the 3D coordinates (x, y, z) . The signed distance field \mathcal{F} becomes $\mathcal{F}(\mathbf{p}, t, \mathbf{c}_i) = \mathcal{T}(\mathcal{D}_1(\mathbf{p}, \mathbf{c}_{ip}), \mathcal{D}_2(t, \mathbf{c}_{it})) = \mathcal{T}(\mathbf{p}', t') = s$, where $s \in \mathbb{R}$ is the signed distance to the closest surface, $\mathcal{T} : \mathbb{R}^4 \rightarrow \mathbb{R}$ is the learned implicit atlas parameterised as a single-shape DeepSDF (Park et al., 2019), $\mathcal{D}_1 : \mathbb{R}^3 \times \mathbb{R}^m \rightarrow \mathbb{R}^3$ is a spatial diffeomorphic deformation module conditioned on spatial latent code $\mathbf{c}_{ip} \in \mathbb{R}^m$, and $\mathcal{D}_2 : \mathbb{R} \times \mathbb{R}^n \rightarrow \mathbb{R}$ is a temporal diffeomorphic deformation module conditioned on temporal latent code $\mathbf{c}_{it} \in \mathbb{R}^n$. Here \mathbf{p}' and t' denote the warped spatial and temporal coordinates in the template space, and \mathbf{c}_i is decomposed into \mathbf{c}_{ip} and \mathbf{c}_{it} for the spatial and temporal deformations, respectively. The spatiotemporal atlas is learned implicitly during training without explicit supervision.

3. Experiments and Results

Data Preprocessing: We selected 100 subjects from the UK Biobank CMR database (80 training, 20 testing). CMR slices were segmented using nnU-Net (Isensee et al., 2021) and reconstructed into dense 3D volumes per frame via a shape completion pipeline (Xu et al., 2025). Left ventricular meshes were extracted via Marching Cubes (Lorenson and Cline, 1987) and processed following DeepSDF (Park et al., 2019) to sample 3D coordinates and compute signed distance values, with the first-frame bounding box applied across all frames

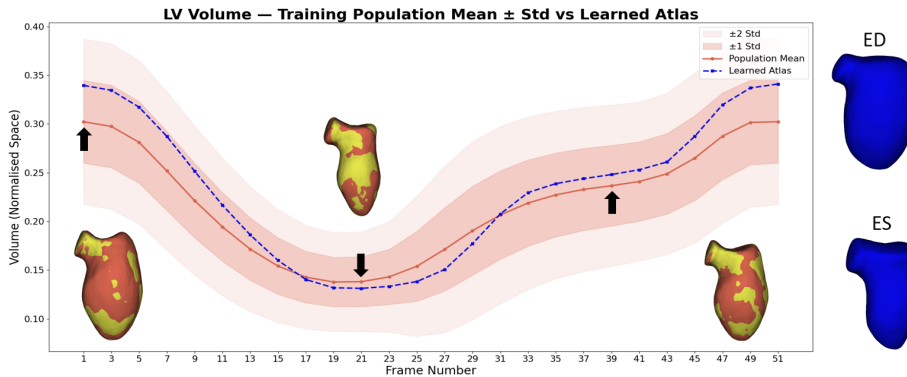


Figure 2: Volume curves of the left ventricle from the learned atlas compared to the training population, computed in the normalised space. ED and ES frames of the learned atlas are shown on the right (blue). Alignment between ground truth (red) and reconstructed meshes from the ED frame (yellow) is presented at different cardiac phases (indicated by black arrows): ED, ES, and diastasis.

to ensure temporal coherence. Each 50-frame sequence was made cyclic by duplicating the first frame as the last, then subsampled to 26 frames, with withheld frames serving as ground truth for motion recovery evaluation. Time coordinates were normalised to $[0, 1]$ and paired with their corresponding sampled 3D coordinates.

Atlas Generation and Motion Recovery Figure 2 shows the left ventricular volume curves computed in the learned template space compared to the training population average. The curves follow the expected contraction–relaxation pattern and differ from the population mean, indicating that the atlas captures meaningful cardiac dynamics. For motion recovery, we report chamfer distance (CD) and 95th percentile Hausdorff distance (HD95) under three settings: (1) 26 frames (same as training), (2) end-diastolic (ED) and end-systolic (ES) frames only, and (3) ED frame only (CT-style). The CD / HD95 results (mm, mean \pm std) are: 1.03 ± 0.25 / 2.40 ± 0.68 (26 frames), 1.24 ± 0.57 / 2.91 ± 1.60 (ED+ES), and 1.36 ± 0.49 / 3.25 ± 1.34 (ED only). Even with a single ED frame, the model achieves comparable accuracy, improving as more frames are provided at test time.

4. Conclusion and Future Work

We extend NDF to spatiotemporal atlas learning, demonstrating strong motion recovery on CMR sequences. Future work includes extending beyond the left ventricle to additional structures and pathological cases, and benchmarking against existing methods.

Acknowledgments

YX and AAY acknowledge funding from EPSRC (EP/S022104/1, Z533762) and Wellcome/EPSRC (WT203148/Z/16/Z). YX acknowledges stipend funding from Heartflow Inc. MS and EP are employees of Heartflow Inc. This research has been conducted using the UK Biobank Resource under application number 88878.

References

- Wenjia Bai, Wenzhe Shi, Antonio De Marvao, Timothy J.W. Dawes, Declan P. O’Regan, Stuart A. Cook, and Daniel Rueckert. A bi-ventricular cardiac atlas built from 1000+ high resolution MR images of healthy subjects and an analysis of shape and motion. *Medical Image Analysis*, 26(1):133–145, December 2015. ISSN 13618415. doi: 10.1016/j.media.2015.08.009. URL <https://linkinghub.elsevier.com/retrieve/pii/S1361841515001346>.
- Fabian Isensee, Paul F. Jaeger, Simon A. A. Kohl, Jens Petersen, and Klaus H. Maier-Hein. nnu-net: a self-configuring method for deep learning-based biomedical image segmentation. *Nature Methods*, 18(2):203–211, 2021. doi: 10.1038/s41592-020-01008-z.
- William E. Lorensen and Harvey E. Cline. Marching cubes: A high resolution 3d surface construction algorithm. *ACM SIGGRAPH Computer Graphics*, 21(4):163–169, 1987. doi: 10.1145/37401.37422.
- Qiang Ma, Qingjie Meng, Mengyun Qiao, Paul M. Matthews, Declan P. O’Regan, and Wenjia Bai. CardiacFlow: 3D+t Four-Chamber Cardiac Shape Completion and Generation via Flow Matching, 2025. URL <https://arxiv.org/abs/2509.05754>. Version Number: 1.
- Shanthi Mendis, Pekka Puska, and Bo Norrving, editors. *Global atlas on cardiovascular disease prevention and control*. World Health Organization, Geneva, 2011. ISBN 9789241564373. URL <https://www.who.int/publications/i/item/9789241564373>. Published in collaboration with the World Heart Federation and the World Stroke Organization.
- Jeong Joon Park, Peter Florence, Julian Straub, Richard Newcombe, and Steven Lovegrove. DeepSDF: Learning continuous signed distance functions for shape representation. In *Proceedings of the IEEE/CVF Conference on Computer Vision and Pattern Recognition (CVPR)*, pages 165–174, 2019. doi: 10.1109/CVPR.2019.00025. URL https://openaccess.thecvf.com/content_CVPR_2019/html/Park_DeepSDF_Learning_Continuous_Signed_Distance_Functions_for_Shape_Representation_CVPR_2019_paper.html.
- Steffen E. Petersen, Paul M. Matthews, Jane M. Francis, Matthew D. Robson, Filip Zemrak, Redha Boubertakh, Alistair A. Young, Sarah Hudson, Peter Weale, Steve Garratt, Rory Collins, Stefan Piechnik, and Stefan Neubauer. UK Biobank’s cardiovascular magnetic resonance protocol. *Journal of Cardiovascular Magnetic Resonance*, 18(1):8, January 2016. ISSN 10976647. doi: 10.1186/s12968-016-0227-4. URL <https://linkinghub.elsevier.com/retrieve/pii/S1097664723009493>.
- Mengyun Qiao, Shuo Wang, Huaqi Qiu, Antonio De Marvao, Declan P. O’Regan, Daniel Rueckert, and Wenjia Bai. CHeart: A Conditional Spatio-Temporal Generative Model for Cardiac Anatomy. *IEEE Transactions on Medical Imaging*, 43(3):1259–1269, March 2024. ISSN 0278-0062, 1558-254X. doi: 10.1109/TMI.2023.3331982. URL <https://ieeexplore.ieee.org/document/10315018/>.

Cathie Sudlow, John Gallacher, Naomi Allen, Valerie Beral, Paul Burton, John Danesh, Paul Downey, Paul Elliott, Jane Green, Martin Landray, Bette Liu, Paul Matthews, Gina Ong, Jill Pell, Alan Silman, Alan Young, Tim Sprosen, Tim Peakman, and Rory Collins. Uk biobank: an open access resource for identifying the causes of a wide range of complex diseases of middle and old age. *PLoS Medicine*, 12(3): e1001779, 2015. doi: 10.1371/journal.pmed.1001779. URL <https://journals.plos.org/plosmedicine/article?id=10.1371/journal.pmed.1001779>.

Shanlin Sun, Kun Han, Deying Kong, Hao Tang, Xiangyi Yan, and Xiaohui Xie. Topology-Preserving Shape Reconstruction and Registration via Neural Diffeomorphic Flow. In *2022 IEEE/CVF Conference on Computer Vision and Pattern Recognition (CVPR)*, pages 20813–20823, New Orleans, LA, USA, June 2022. IEEE. ISBN 978-1-6654-6946-3. doi: 10.1109/CVPR52688.2022.02018. URL <https://ieeexplore.ieee.org/document/9880332/>.

Yiyang Xu, Hao Xu, Matthew Sinclair, Esther Puyol-Antón, Steven A. Niederer, Amedeo Chiribiri, Steven E. Williams, Michelle C. Williams, and Alistair A. Young. Improved 3d whole heart geometry from sparse cmr slices. In Oscar Camara, Esther Puyol-Antón, Maxime Sermesant, Avan Suinesiaputra, Jichao Zhao, Chengyan Wang, Qian Tao, and Alistair Young, editors, *Statistical Atlases and Computational Models of the Heart. Workshop, CMRxRecon and MBAS Challenge Papers.*, pages 43–52, Cham, 2025. Springer Nature Switzerland. ISBN 978-3-031-87756-8.

Alistair A. Young and Alejandro F. Frangi. Computational cardiac atlases: from patient to population and back. *Experimental Physiology*, 94(5):578–596, May 2009. ISSN 0958-0670, 1469-445X. doi: 10.1113/expphysiol.2008.044081. URL <https://physoc.onlinelibrary.wiley.com/doi/10.1113/expphysiol.2008.044081>.