

# Self-Supervised Cortical Surface Reconstruction for Ultra High-resolution *ex vivo* 7T MRI

Haoxiang Li<sup>\*1</sup>

Mingxuan Liu<sup>\*1</sup>

Hongjia Yang<sup>1</sup>

Yi Liao<sup>2</sup>

Haibo Qu<sup>2</sup>

Jonathan Polimeni<sup>3</sup>

Qiyuan Tian<sup>†1</sup>

LIHAOXIA24@MAILS.TSINGHUA.EDU.CN

ARKTISX@FOXMAIL.COM

YANGHJ23@MAILS.TSINGHUA.EDU.CN

CONNIE0064@126.COM

WINDOWSQHB@126.COM

JRPOLIMENI@STANFORD.EDU

QIYUANTIAN@TSINGHUA.EDU.CN

<sup>1</sup> School of Biomedical Engineering, Tsinghua University

<sup>2</sup> Department of Radiology, West China Second University Hospital, Sichuan University

<sup>2</sup> Department of Radiology, Stanford University

**Editors:** Under Review for MIDL 2025

## Abstract

*Ex vivo* brain MRI enables sub-millimeter ultra-high-resolution studies, uncovering structural details unattainable with *in vivo* MRI. Cortical surface reconstruction (CSR) based on these detailed images is crucial for studying cortical anatomy and structure. Despite this potential, methodological development in *ex vivo* MRI has been constrained by several factors: scarcity of datasets, limited imaging resources, pronounced susceptibility artifacts, and signal inhomogeneity. While learning-based CSR methods have been proposed to accelerate reconstruction processes, they face a fundamental limitation—requiring CSR results from classic methods like FreeSurfer as training references, making them mostly only working for *in vivo* adult MRI data and unsuitable for the unique characteristics of *ex vivo* brain imaging. To address the challenge, we propose **SelfCSR**, a self-supervised deep learning framework for accurate *ex vivo* 7T MRI CSR without the need for manually labeled training data.

**Keywords:** *Ex vivo* Brain, 7T MRI, Self-supervised Learning, Cortical Surface Reconstruction.

## 1. Introduction

*Ex vivo* MRI offers significant advantages over *in vivo* MRI by enabling detailed neuroanatomy visualization, bridging microscale histology studies with morphometric measurements, and linking macroscopic features such as cortical thickness to underlying cytoarchitecture and pathology (Khandelwal et al., 2024). Meanwhile, compared to standard 1.5T or 3T MRI, 7T MRI provides ultra-high resolution and significantly enhanced contrast, making it an invaluable tool for detailed neuroimaging. Since *ex vivo* MRI can be conducted with less time constraints (e.g., lasting days) and is free from cardiorespiratory or head motion, it enables the application of 7T imaging for brain neuroscientific studies.

---

\* Contributed equally

† Corresponding author

For example, Coras et al. (Coras et al., 2014) leveraged 7T *ex vivo* MRI to characterize the microstructural differences between normal and sclerotic hippocampi in temporal lobe epilepsy. Zeng et al. (Zeng et al., 2024) developed a segmentation model specifically for 7T *ex vivo* MRI data using manually labeled training datasets. Another study also developed a deep learning pipeline for automated high-resolution postmortem MRI segmentation, linking cortical and subcortical morphometry to neuropathology in neurodegenerative diseases (Khandelwal et al., 2024).

Cortical surface reconstruction (CSR) for *ex vivo* brain is crucial for studying cortical anatomy and structure. However, existing CSR tools, such as FreeSurfer (Fischl, 2012) and BrainSuite (Shattuck and Leahy, 2002), struggle to process 7T MRI *ex vivo* brain data due to its unique tissue contrast and ultra-high resolution. Deep learning provides a promising approach for CSR (Ma et al., 2022; Li et al., 2025). However, due to the scarcity of datasets, limited imaging resources, pronounced susceptibility artifacts, and signal inhomogeneity, existing supervised learning methods like CortexODE (Ma et al., 2022) and CoTAN (Ma et al., 2023) are not applicable to *ex vivo* brain imaging. Recent advances, such as SegCSR (Zheng et al., 2024) and CoSeg (Ma et al., 2024), have explored self-supervised CSR for *in vivo* T1-weighted brain MRI by leveraging pseudo ground truth synthesized from segmentation. Despite these advancements, there remains a lack of methods tailored for cortical surface reconstruction directly from 7T MRI data. To address this gap, we present **SelfCSR**, the first self-supervised CSR method tailored for 7T *ex vivo* brain MRI. **SelfCSR** harnesses the unique advantages of 7T MRI while addressing challenges from its distinct tissue contrast and high resolution, enabling more accurate and efficient cortical surface analysis.

## 2. Method

### 2.1. Data Description

This study used the publicly available 7 Tesla *ex vivo* human brain MRI dataset at 100-micron resolution from Edlow et al. (Edlow et al., 2019). The scans were acquired using a 7 Tesla whole-body human MRI scanner with four single-echo spoiled gradient-recalled echo (SPGR/GRE) or Fast Low-Angle Shot (FLASH) sequences. The dataset features exceptionally high-resolution images (1600×1400×640 voxels), with each dataset approximating 4.9GB in size.

### 2.2. *ex vivo* Brain Cortex Surface Reconstruction Pipeline

The proposed **SelfCSR** consists of four stages: automatic segmentation, downsampling and denoising, marching cubes, and self-supervised surface deformation.

**Automatic Segmentation.** The segmentation of supragranular and infragranular layers is achieved using the semi-supervised multi-resolution U-Net framework proposed by Zeng et al (Zeng et al., 2024).

**Downsampling and Denoising.** The data is first downsampled to an isotropic resolution of 0.5 mm, significantly reducing memory consumption while preserving essential anatomical details. Furthermore, the BM4D algorithm (Maggioni et al., 2012) is applied for denoising to enhance data quality.

**Marching Cube.** The marching cubes algorithm (WE, 1987) is employed to extract three cortical surfaces from the segmentation: the white matter (WM) surface, the mid-surface (representing the boundary between the supragranular and infragranular layers), and the pial surface. These surfaces collectively provide a comprehensive representation of cortical structures. The WM surface is used as the initial input and reference geometry for subsequent deformation, while the mid-surface and pial surface are used as pseudo-labels for training in the framework.

**Surface Deformation.** To ensure point-wise correspondence between the three generated surfaces, we employ two U-Net models to predict stationary velocity fields (SVFs). The first U-Net deforms the white matter surface to align with the mid-surface, and the second U-Net subsequently deforms the mid-surface to align with the pial surface. This step establishes precise point-wise relationships across the three surfaces, enabling accurate and consistent cortical surface analysis. The following self-supervised loss functions were employed during training:

$$\mathcal{L} = w_{cd}\mathcal{L}_{cd} + w_{gr}\mathcal{L}_{gr} + w_{it}\mathcal{L}_{it} + w_{edge}\mathcal{L}_{edge} + w_{normal}\mathcal{L}_{normal} \quad (1)$$

The boundary alignment loss ( $\mathcal{L}_{cd}$ ) optimizes surface-to-boundary fit using one-way Chamfer distance, while the gradient loss ( $\mathcal{L}_{gr}$ ) maximizes MRI gradients at the surface for sharper boundaries. The intensity loss ( $\mathcal{L}_{it}$ ) constrains voxel values to plausible ranges. Topological integrity is enforced by a mesh regularity term ( $\mathcal{L}_{edge}$ ) penalizing irregular edge lengths and an inflation constraint ( $\mathcal{L}_{deform}$ ) ensuring normal-direction expansion. The terms  $w_{cd}$ ,  $w_{gr}$ ,  $w_{it}$ ,  $w_{edge}$ , and  $w_{normal}$  serve as weight hyperparameters to balance the contribution of each loss component. Training is performed through an ODE-based framework for efficient optimization.

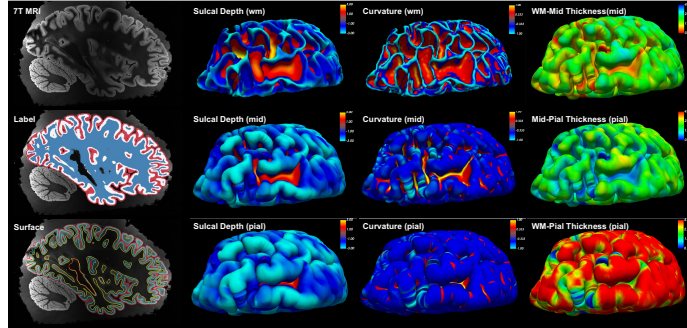


Figure 1: Cortical surface reconstruction and morphological analysis.

### 3. Results and Conclusion

Figure 1 presents the comprehensive cortical reconstruction results, including the segmented white matter (WM) boundary, reconstructed mid-surface, and pial surface. We further analyzed *ex vivo* brain morphology using FreeSurfer (Fischl, 2012) to quantify cortical curvature, thickness, and sulcal depth. In conclusion, **SelfCSR** is the first learning-based *ex vivo* CSR method and offering a novel tool for *ex vivo* MRI-based neuroimaging research.

## Acknowledgments

Tsinghua University Startup Fund and Dushi Program (grant number 20241080026).

## References

- Roland Coras, Gloria Milesi, Ileana Zucca, Alfonso Mastropietro, Alessandro Scotti, Matteo Figini, Angelika Mühlebner, Andreas Hess, Wolfgang Graf, Giovanni Tringali, et al. 7 t mri features in control human hippocampus and hippocampal sclerosis: an ex vivo study with histologic correlations. *Epilepsia*, 55(12):2003–2016, 2014.
- Brian L Edlow, Azma Mareyam, Andreas Horn, Jonathan R Polimeni, Thomas Witzel, M Dylan Tisdall, Jean C Augustinack, Jason P Stockmann, Bram R Diamond, Allison Stevens, et al. 7 tesla mri of the ex vivo human brain at 100 micron resolution. *Scientific data*, 6(1):244, 2019.
- Bruce Fischl. Freesurfer. *Neuroimage*, 62(2):774–781, 2012.
- Pulkit Khandelwal, Michael Tran Duong, Shokufeh Sadaghiani, Sydney Lim, Amanda E. Denning, Eunice Chung, Sadhana Ravikumar, Sanaz Arezoumandan, Claire Peterson, Madigan Bedard, Noah Capp, Ranjit Ittyerah, Elyse Migdal, Grace Choi, Emily Kopp, Bridget Loja, Eusha Hasan, Jiacheng Li, Alejandra Bahena, Karthik Prabhakaran, Gabor Mizsei, Marianna Gabrielyan, Theresa Schuck, Winifred Trotman, John Robinson, Daniel T. Ohm, Edward B. Lee, John Q. Trojanowski, Corey McMillan, Murray Grossman, David J. Irwin, John A. Detre, M. Dylan Tisdall, Sandhitsu R. Das, Laura E. M. Wisse, David A. Wolk, and Paul A. Yushkevich. Automated deep learning segmentation of high-resolution 7 tesla postmortem mri for quantitative analysis of structure-pathology correlations in neurodegenerative diseases. *Imaging Neuroscience*, 2:1–30, 05 2024. ISSN 2837-6056. doi: 10.1162/imag\_a.00171. URL [https://doi.org/10.1162/imag\\_a.00171](https://doi.org/10.1162/imag_a.00171).
- Haoxiang Li, Mingxuan Liu, Xuguang Bai, Yi Liao, Jialan Zheng, Hongjia Yang, Zihan Li, Haibo Qu, and Qiyuan Tian. FetalCSR: Multi-input attention fusion network for neural ODE-based fetal cortical surface reconstruction. In *AI for Children: Healthcare, Psychology, Education*, 2025. URL <https://openreview.net/forum?id=Ra0xioC3He>.
- Qiang Ma, Liu Li, Emma C Robinson, Bernhard Kainz, Daniel Rueckert, and Amir Alansary. Cortexode: Learning cortical surface reconstruction by neural odes. *IEEE Transactions on Medical Imaging*, 42(2):430–443, 2022.
- Qiang Ma, Liu Li, Vanessa Kyriakopoulou, Joseph V Hajnal, Emma C Robinson, Bernhard Kainz, and Daniel Rueckert. Conditional temporal attention networks for neonatal cortical surface reconstruction. In *International Conference on Medical Image Computing and Computer-Assisted Intervention*, pages 312–322. Springer, 2023.
- Qiang Ma, Liu Li, Emma C Robinson, Bernhard Kainz, and Daniel Rueckert. Weakly supervised learning of cortical surface reconstruction from segmentations. In *International Conference on Medical Image Computing and Computer-Assisted Intervention*, pages 766–777. Springer, 2024.

- Matteo Maggioni, Vladimir Katkovnik, Karen Egiazarian, and Alessandro Foi. Nonlocal transform-domain filter for volumetric data denoising and reconstruction. *IEEE transactions on image processing*, 22(1):119–133, 2012.
- David W Shattuck and Richard M Leahy. Brainsuite: an automated cortical surface identification tool. *Medical image analysis*, 6(2):129–142, 2002.
- LORENSEN WE. Marching cubes: A high resolution 3d surface construction algorithm. *Computer graphics*, 21(1):7–12, 1987.
- Xiangrui Zeng, Oula Puonti, Areej Sayeed, Rogeny Herisse, Jocelyn Mora, Kathryn Evancic, Divya Varadarajan, Yael Balbastre, Irene Costantini, Marina Scardigli, et al. Segmentation of supragranular and infragranular layers in ultra-high-resolution 7t ex vivo mri of the human cerebral cortex. *Cerebral Cortex*, 34(9):bhae362, 2024.
- Hao Zheng, Xiaoyang Chen, Hongming Li, Tingting Chen, Peixian Liang, and Yong Fan. Segcsr: Weakly-supervised cortical surfaces reconstruction from brain ribbon segmentations. *bioRxiv*, pages 2024–12, 2024.