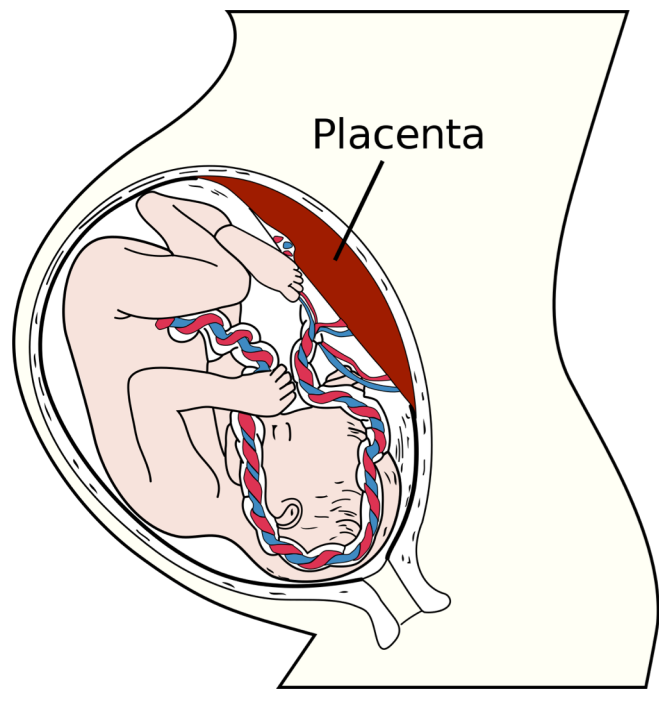


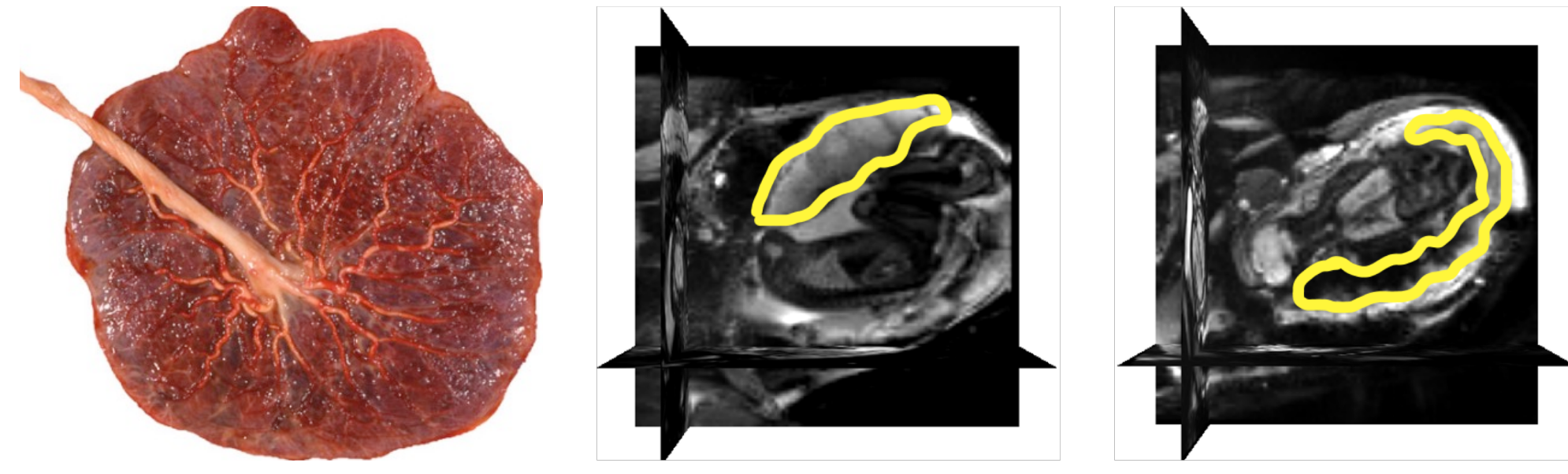
Goal

Improve visualization of placenta function, anatomy



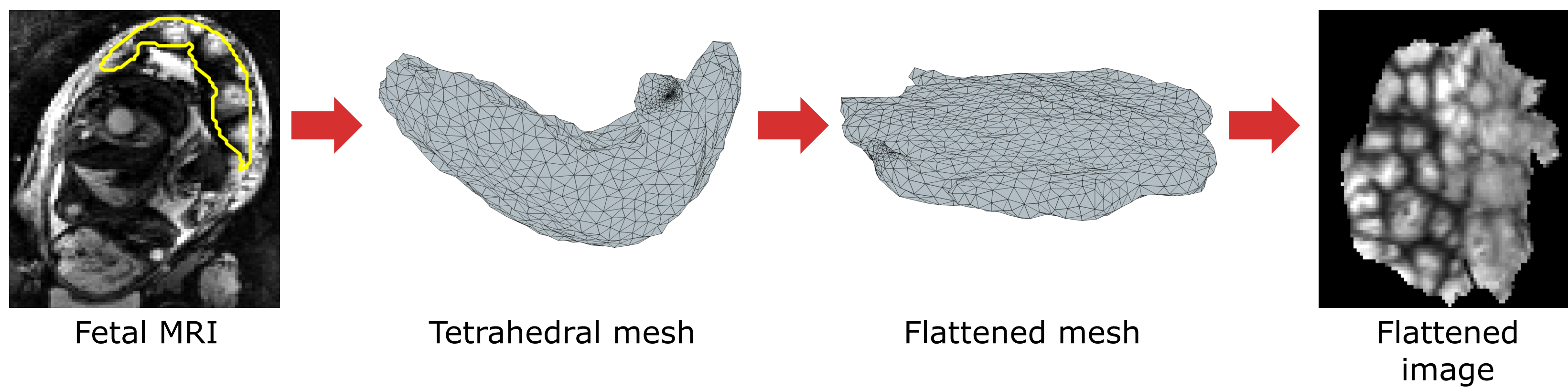
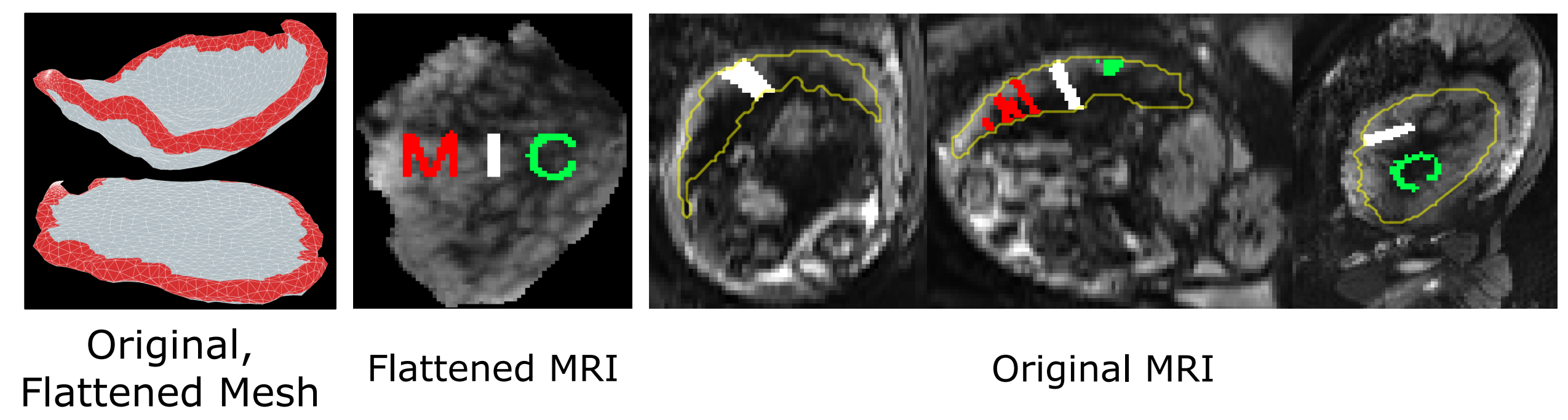
Clinical Motivations

- Prevent pregnancy complications: assess placental function by fetal MRI
- Difficult to visualize, study function: variable and curved placental shape

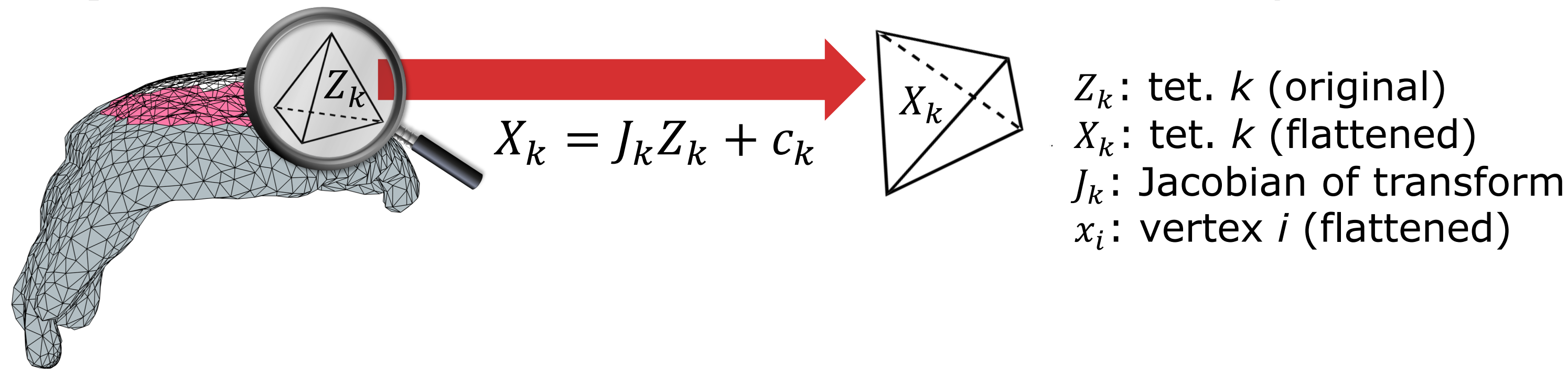


Approach

Map the placenta volume to a flattened template to resemble the well-studied post-birth shape.



1. Map Formulation: Piecewise affine to a template



2. Mapping Objective Function

$$\phi(X, h) = \underbrace{\sum_{m: x_m \in \partial Z} Area_m T(x_m, h)}_{\text{Template Match } (T)} + \lambda \underbrace{\sum_{k=1}^K Vol_k \mathcal{D}(J(X_k))}_{\text{Volumetric Distortion } (D)}$$

- Distortion: Symmetric Dirichlet Energy

$$\mathcal{D}(J) = \|J\|_F^2 + \|J^{-1}\|_F^2 = \sum_{i=1}^3 (\sigma_i^2 + \sigma_i^{-2})$$

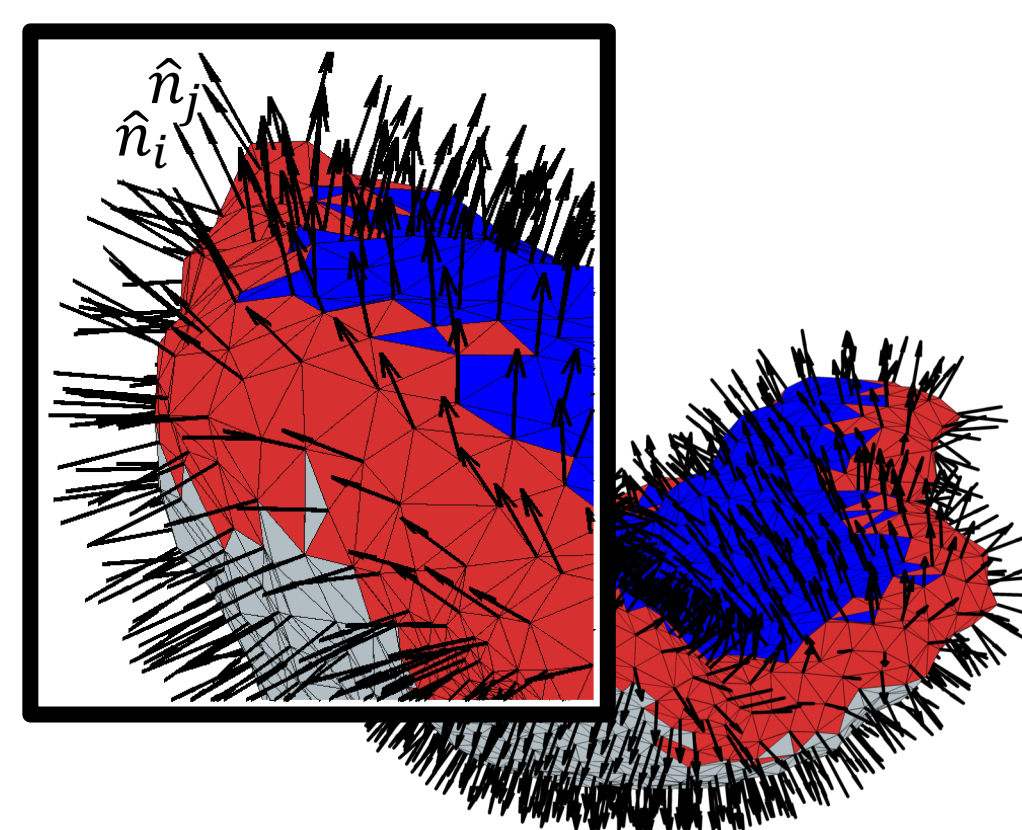
- Template: Uniform thickness

$$T(x, h) = \begin{cases} (x^{(3)} - h)^2, & \text{if } x \in \mathcal{F}(\partial Z) \\ (x^{(3)} + h)^2, & \text{if } x \in \mathcal{M}(\partial Z) \\ 0 & \text{otherwise} \end{cases}$$

- Boundary parcellation: Spectral clustering to find boundary vertex labelling l that minimizes

$$E(l) = \sum_{ij} w_{ij} (l_i - l_j)^2;$$

subject to $\|l\|_2^2 = 1, \sum_i l_i = 0,$
where, $w_{i,j} = \exp\{\gamma(\hat{n}_i^T \hat{n}_j)\}$



3. Optimization

- Gradient descent over vertex locations with line search
- Invertible map: Prevent tet. volume sign change per iteration

$$X_k^{(n+1)} = X_k^{(n)} - \eta \nabla_X \phi(X)$$

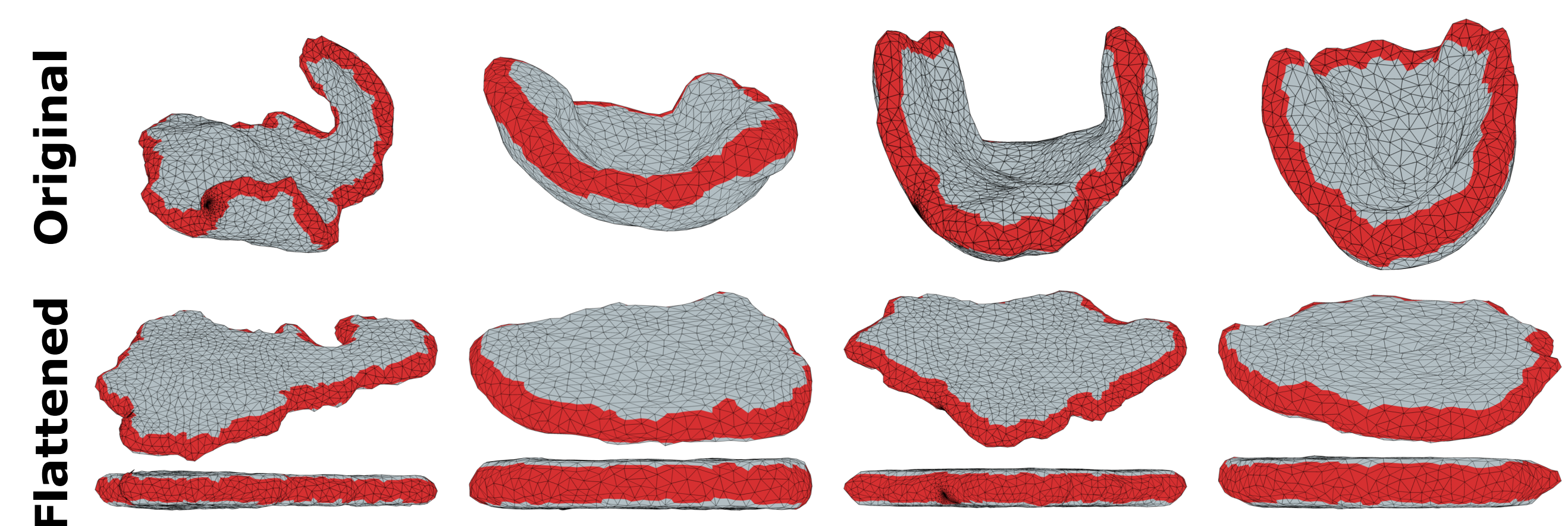
$$Vol_k(\eta) = \frac{1}{6} \det((X_k - \eta \nabla_X \phi(X_k))B)$$

Dataset:

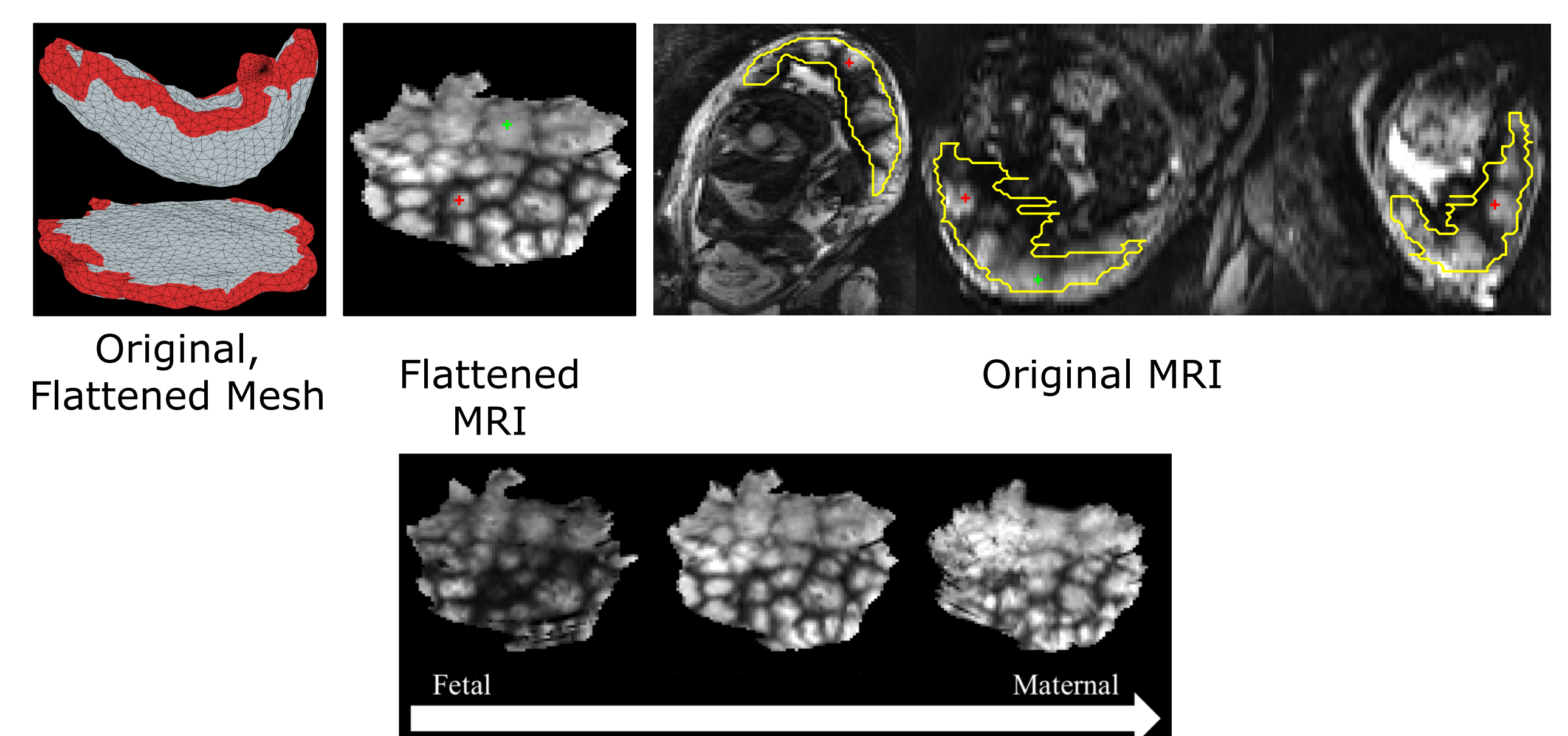
- 78 subjects, gestational age: 27-38 weeks
- 111 segmentations; 59 singleton (35 in supine and lateral positions), 17 twin pregnancies
- MRI: GRE-EPI 3mm isotropic

Results:

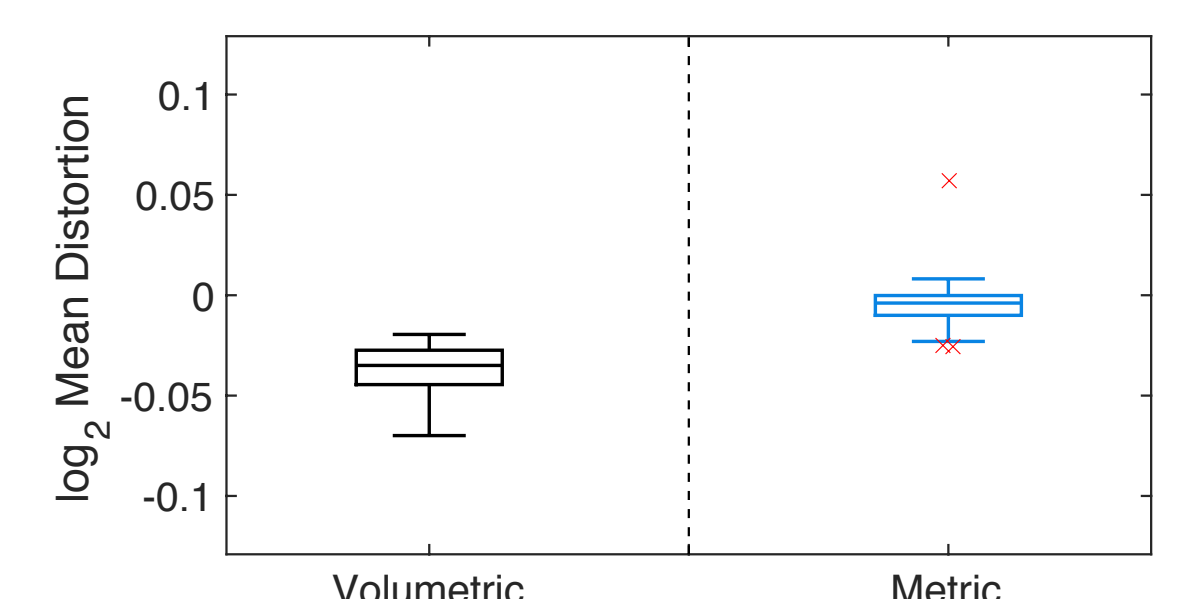
- Mapping is robust to shape variation



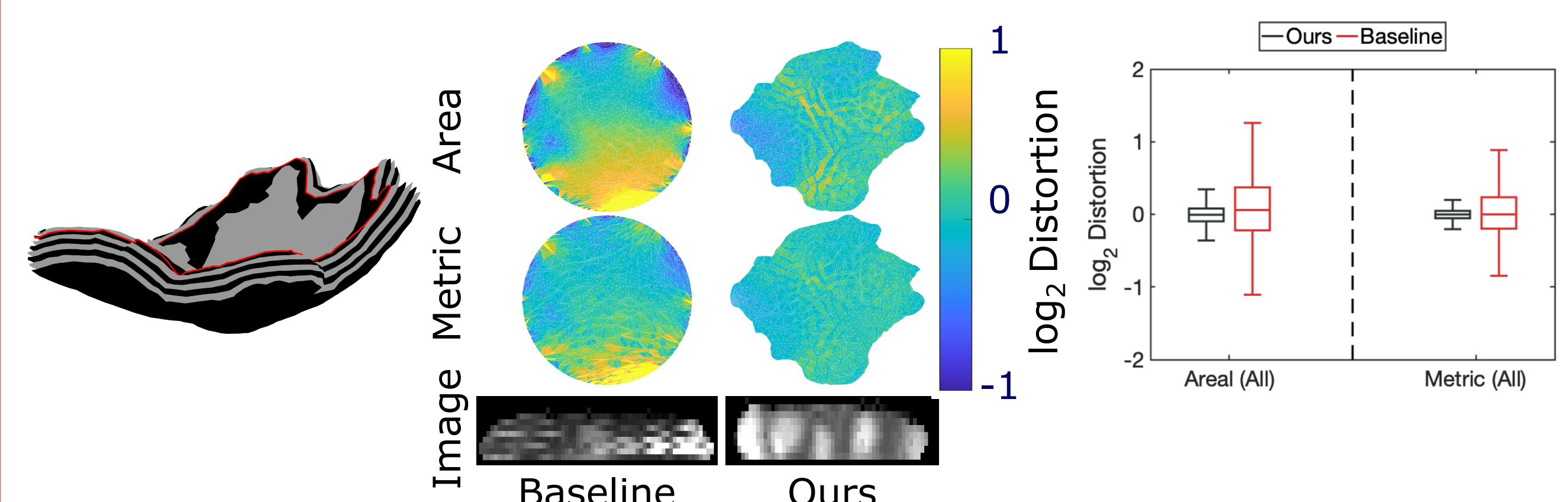
- Enables Contextual Visualization



- Shape distortion: $4.1 \pm 1.9\%$
- Template mismatch 0.26 ± 0.05 voxels



- Improved distortion, visualization over 2D baseline¹



- Next Steps:** *ex vivo* analysis and comparison

