# Non-rigid Registration with Missing Correspondences in Preoperative and Postresection Brain Images

Nicha Chitphakdithai and James S. Duncan Image Processing and Analysis Group, Yale University

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#### **Image Registration Goal**

 Align postresection and preoperative brain MRI of epilepsy patients



- Challenge: Missing Correspondences
  - Cause misalignment of other actual corresponding features

# Approaches to Handle Missing Correspondences

#### Previous Methods

- Adapted demons registration + level set segmentation of resection<sup>1</sup>
- "De-enhance" DCE-MRI before registration<sup>3</sup>
- Estimate registration and missing data assuming equal chance of missing/valid label<sup>4</sup>
- Spatial prior on valid tissue/resection locations for post-resection images<sup>5</sup>

#### Our Approach

- → Jointly register and classify correspondence regions in statistical parameter estimation framework<sup>2</sup>
- → Put less weight on voxels believed to be missing correspondence
- → Include intensity prior on resection voxels

<sup>1</sup>Risholm et al., IPMI 2009; <sup>2</sup>Pohl et al., Neuroimage 2006; <sup>3</sup>Zheng et al., MICCAI 2007; <sup>4</sup>Periaswamy and Farid, MedIA 2006; <sup>5</sup>Chitphakdithai and Duncan, ISBI 2010

# Registration and Indicator Map Estimation (RIME): Overview

• Introduce "hidden" indicator map to segment valid tissue, resection, and background



- Given indicator map
  - → easier registration problem
- Given correct alignment
  → easier to classify regions
- Maximum a posteriori framework:

 $\hat{T} = \arg\max_{T} \log\sum_{I} p(T, I \mid P, R)$ 



# Registration and Indicator Map Estimation (RIME): EM Algorithm

#### • E-Step: Indicator Map Estimation

# • M-Step: Registration E-Step Weights Similarity Term $T^{k+1} = \arg \max_{T} \sum_{\mathbf{x} \in R} \sum_{l \in L} p(I(\mathbf{x}) = l | P, R, T^{k}) \left[ \log p(P(T(\mathbf{x})) | R, I(\mathbf{x}) = l, T) + \log p(R(\mathbf{x}) | I(\mathbf{x}) = l) \right] + \log p(T)$ $+ \log p(R(\mathbf{x}) | I(\mathbf{x}) = l) + \log p(T)$ Intensity Prior Transformation Prior

# Probability Models: Likelihood

- Likelihood  $p(P(T(\mathbf{x}))|R,I(\mathbf{x})=l,T)$  acts like the similarity metric
- Key: For different indicator values, can use different probability models

$$P(T(\mathbf{x})) | R, T, I(\mathbf{x}) = l \sim \begin{cases} Unif(\frac{1}{c}) & ,l = \text{resection} \\ N(R(\mathbf{x}), \sigma_1) & ,l = \text{valid tissue} \\ N(R(\mathbf{x}), \sigma_2) & ,l = \text{background} \end{cases}$$

• Choose c = number of intensity levels  $\sigma_2 > \sigma_1$ 

# Probability Models: Intensity Prior

•  $p(R(\mathbf{x})|I(\mathbf{x})=l)$  incorporates prior knowledge of intensities in postresection image

$$R(\mathbf{x}) | I(\mathbf{x}) = l \sim \begin{cases} N(\mu_r, \sigma_r) &, l = \text{resection} \\ Unif(\frac{1}{c}) &, l = \text{valid tissue} \\ N(0, \sigma_b) &, l = \text{background} \end{cases}$$

 Use training set of manually segmented postoperative images to estimate resection class parameters



# Probability Models: Transformation Prior

- Chose free form deformations (FFDs) based on uniform cubic B-splines
- Assume control points  $\mathbf{t}_{\mathbf{i}}$  and components  $t_{\mathbf{i},\mathbf{j}}$  independent with spacing  $\delta$

$$p(T) = \prod_{i} \prod_{j} p(t_{i,j})$$

$$t_{i,j} \sim N\left(\mu_{i,j}, \frac{0.4\delta}{3}\right)$$

• Restrict control points to lie within sphere of radius  $0.4\delta$  for injective transformation<sup>1</sup>



<sup>1</sup>Greene et al., MedIA 2009

#### Registration Methods for Comparison

- "Standard" non-rigid registration (SNRR) using
  - Uniform cubic B-spline FFDs<sup>1</sup>
  - Sum of squared differences (SSD) similarity
- Robust SSD similarity metric (RTR):  $\rho_s \left( R(\mathbf{x}) - P(T(\mathbf{x})) \right) \sim N(0, \sigma)$ ,  $\rho$  is Tukey function

Tukey function with scaling parameter *s* 



<sup>1</sup>Rueckert et al., TMI 1999

# Synthetic Data: Experimental Setup

#### • Synthetic Image Creation:



 Leave-one-out validation: train intensity prior on 10 images

# Synthetic Data: Registration Results



#### Valid Tissue Map



#### Average Displacement Field Errors 0.4 0.3 0.2 0.1 0 Min\*1e2 Max\*1e-1 Mean Std Dev SNRR RTR RTR RIME

Average dice: 0.98

Significant
 compared to SNRR
 Significant
 compared to RTR

#### Real Data: Registration Results



Average Landmark Errors (6 datasets)

2.69 mm

2.16 mm

1.27 mm

#### Real Data: Indicator Map Estimation



Left temporal lobe resection Pink = valid correspondences, Grey = resection

- Average dice coefficient for valid correspondence estimate: 0.92
- Some mislabeling of valid correspondence voxels as resection

#### **Conclusions and Future Work**

#### • Contributions

- Registration handles missing correspondences by incorporating "hidden" indicator map
- Probabilistic framework allowed inclusion of prior on postresection intensity given the label
- Future Work
  - Image histogram-based likelihood model
  - Improve estimate of indicator map by incorporating spatial prior
    - Include knowledge of resection location
    - Smooth map estimate

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- Contact: nicha.chitphakdithai@yale.edu

