



# Joint Tumor Segmentation and Dense Deformable Registration of Brain MR Images

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2. Equipe GALEN, INRIA Saclay-Ile de France

3. Intrasense SAS, Montpellier

4. Département de Neurochirurgie, Hôpital Gui de Chauliac, Montpellier

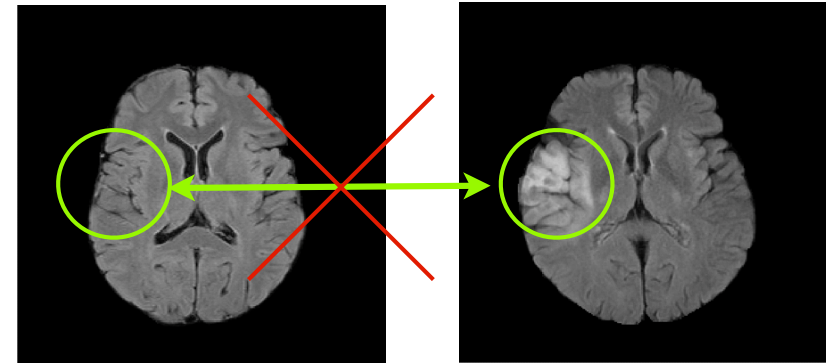
# Introduction

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## Brain Tumor Segmentation and Registration from healthy to pathological subject treated separately



- Fuzzy boundaries
- inhomogeneous appearances
- Various shapes
- intensity overlap with healthy tissue



- No correspondences in the tumor area: use of common methods impossible

### Methods

- Classification techniques + pairwise smoothing  
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- Atlas based segmentation: dependent on registration quality  
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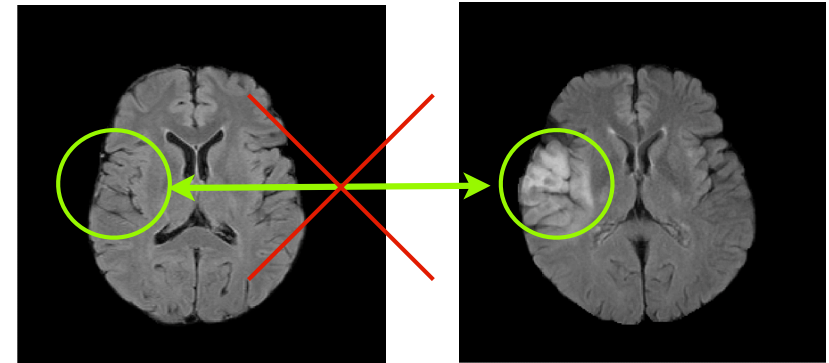
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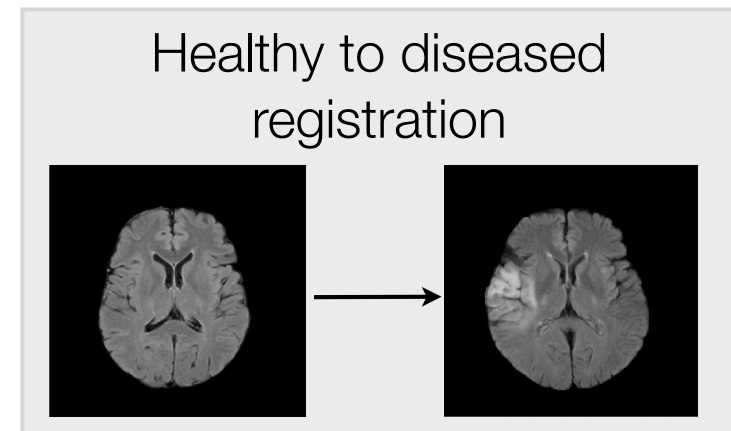
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# Method Overview

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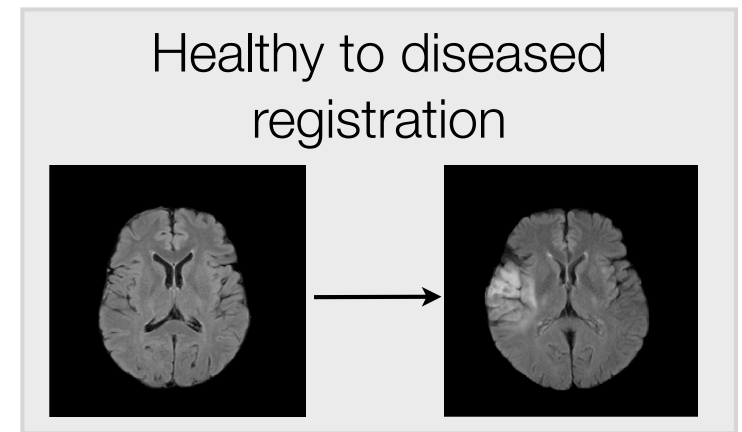
**Simultaneously** register a healthy subject to a diseased subject and find the tumor's segmentation map



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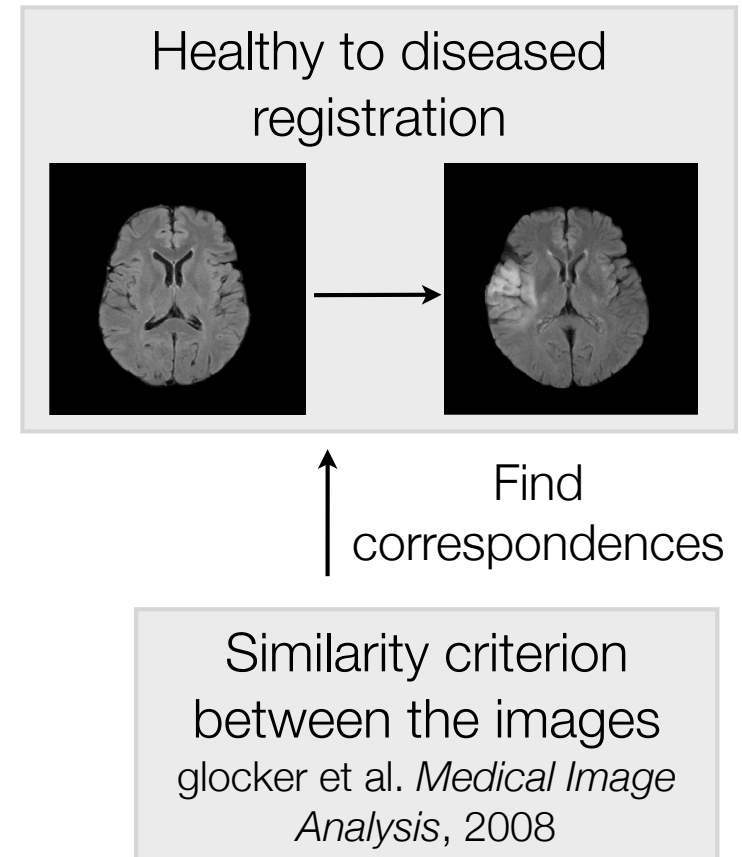
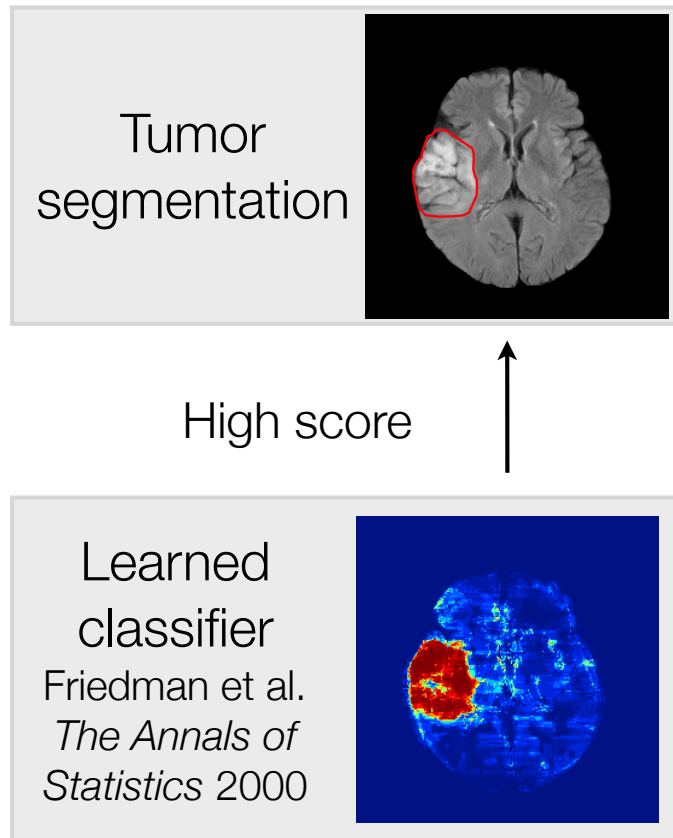


Find correspondences

Similarity criterion  
between the images  
glocker et al. *Medical Image Analysis*, 2008

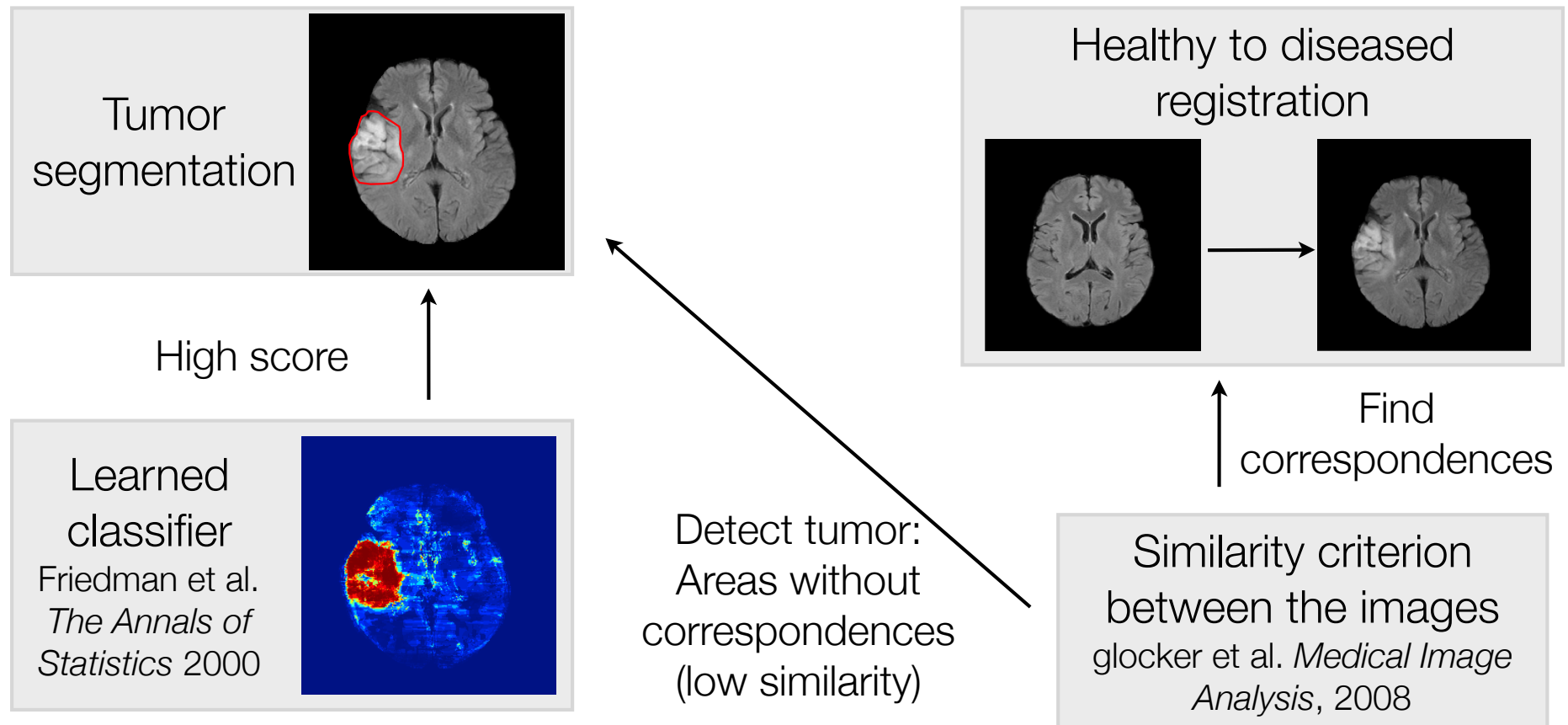
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**Simultaneously** register a healthy subject to a diseased subject and find the tumor's segmentation map



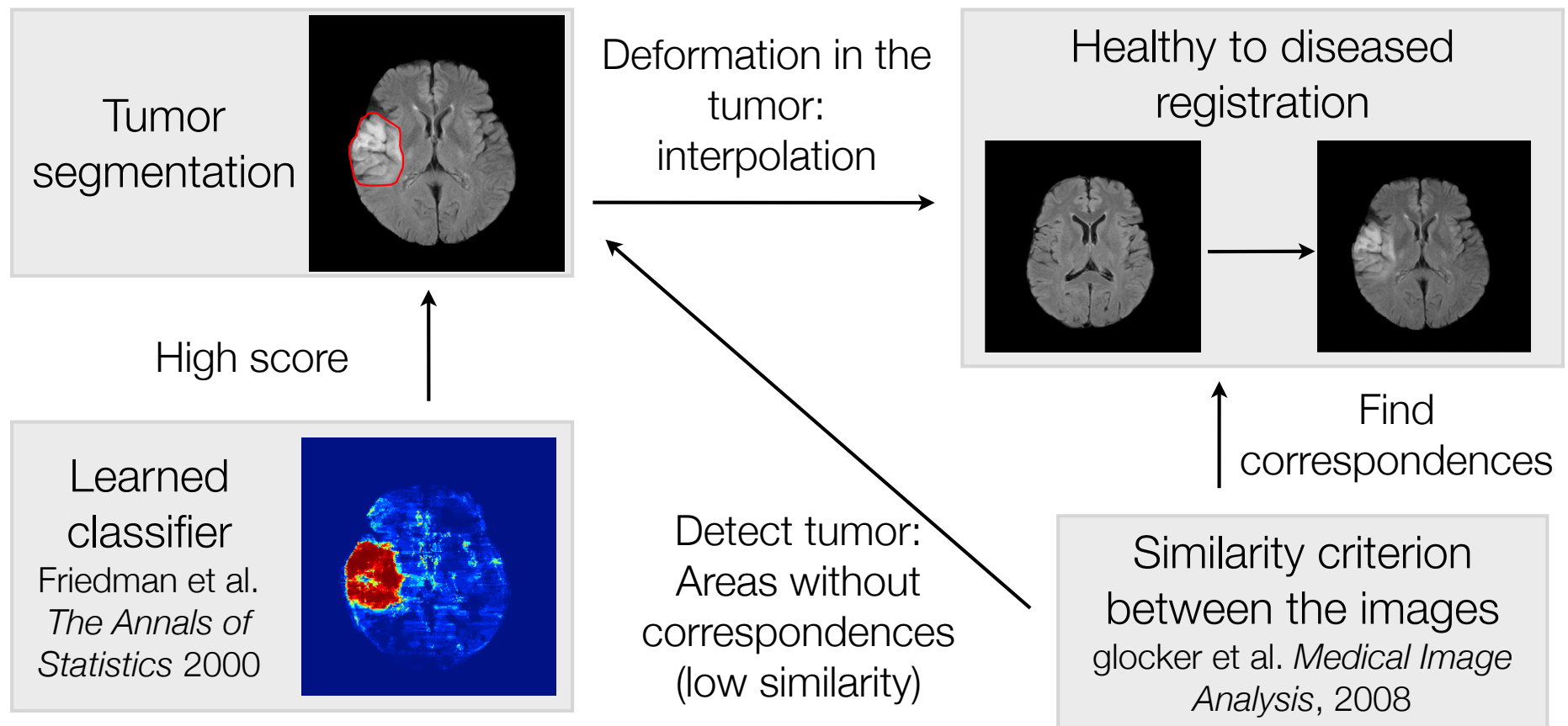
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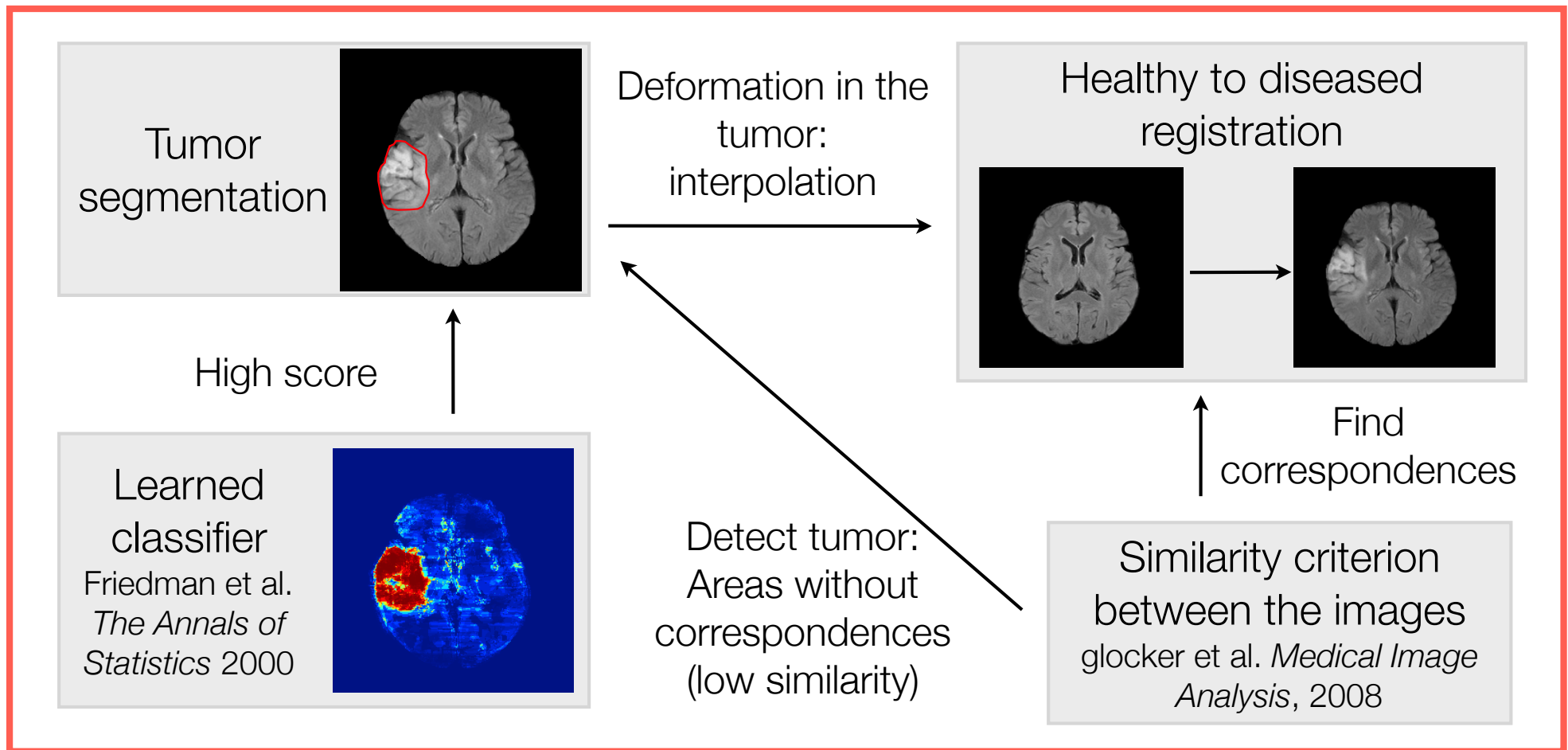
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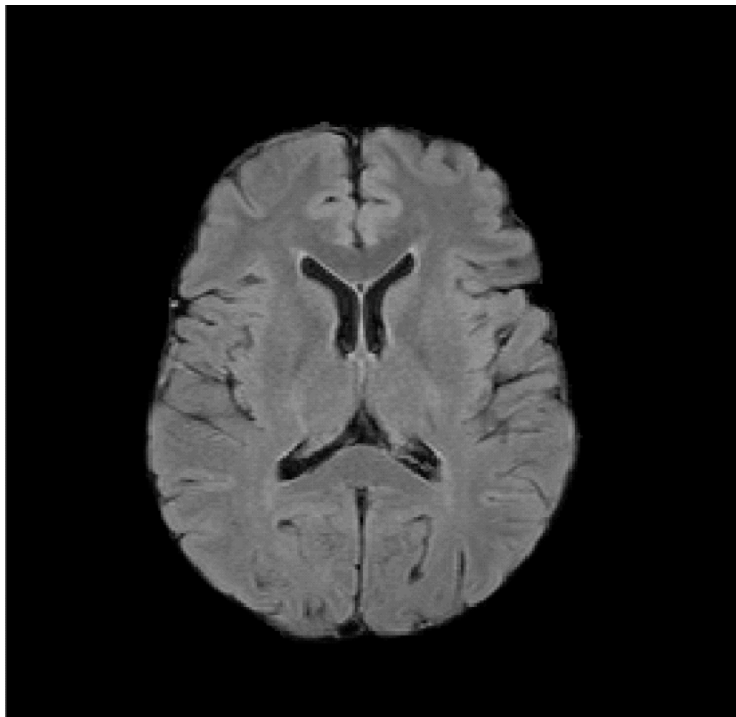


## Discrete Markov Random Field Formulation

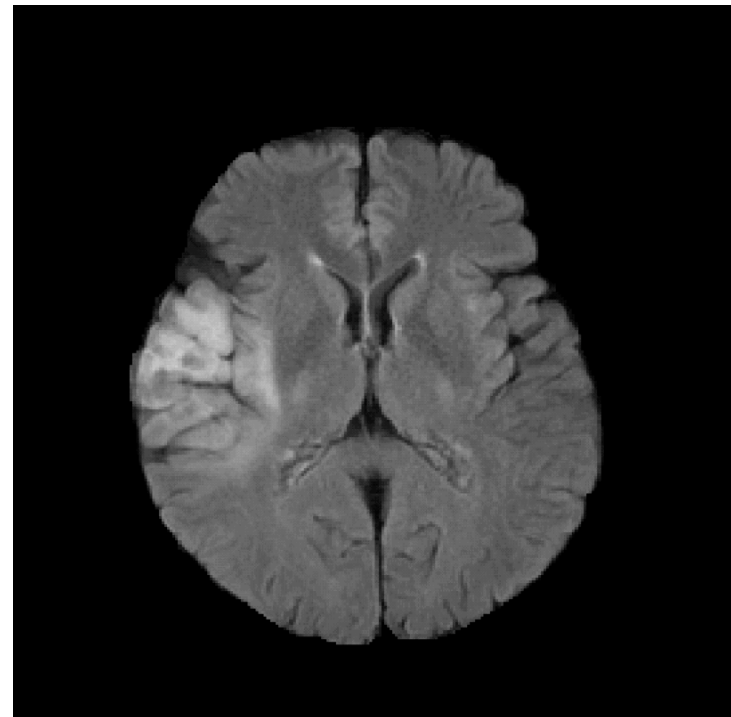
# Parametrization

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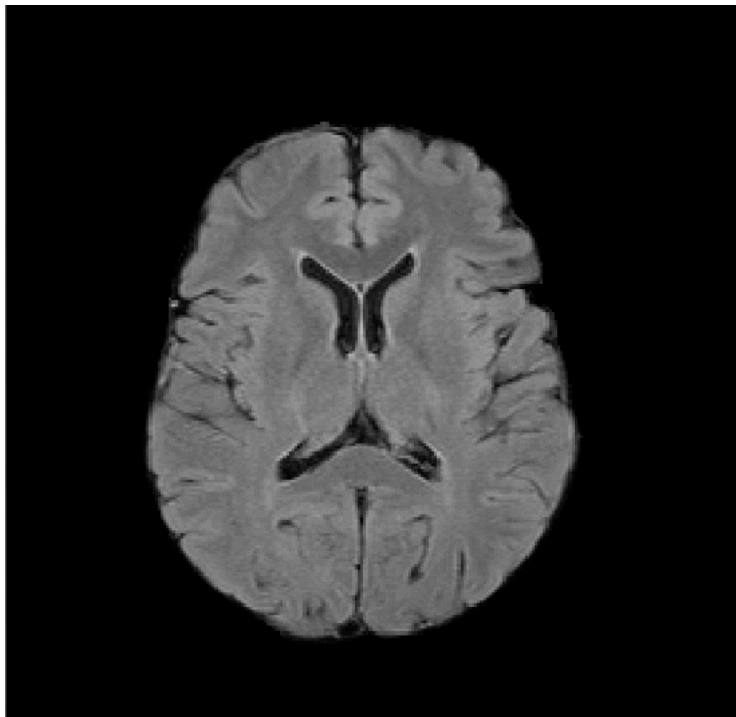


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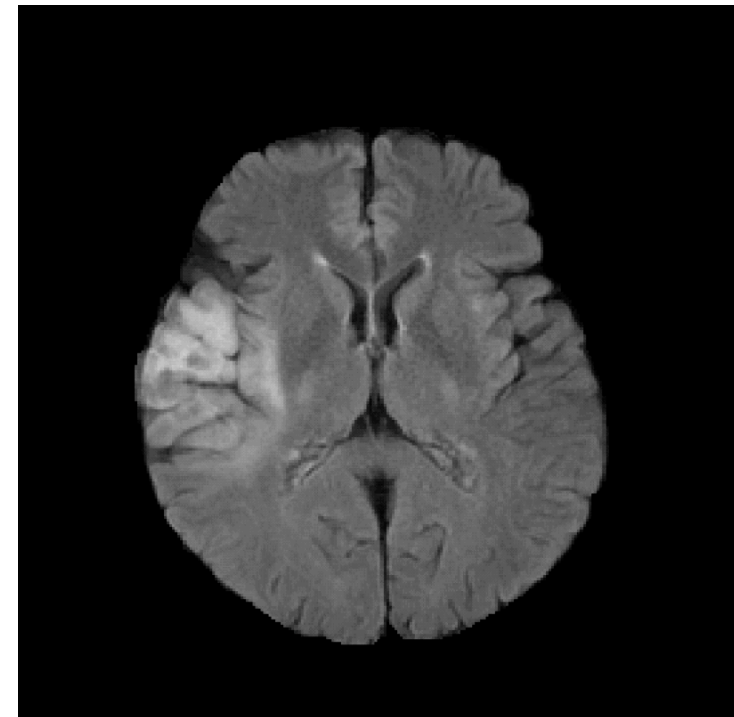
Deformation field

$$\mathcal{T}(\mathbf{x})$$

$$A \circ \mathcal{T}(\mathbf{x}) = I(\mathbf{x})$$

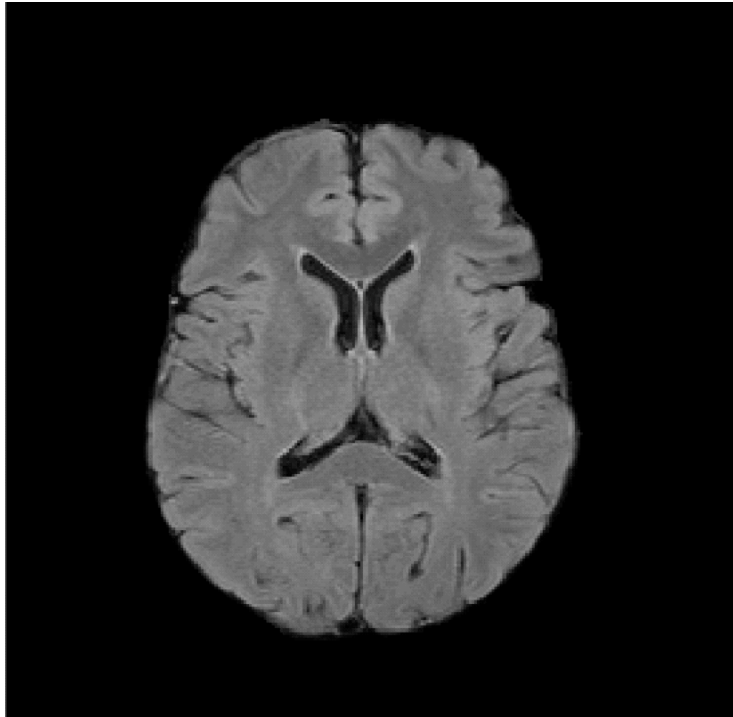


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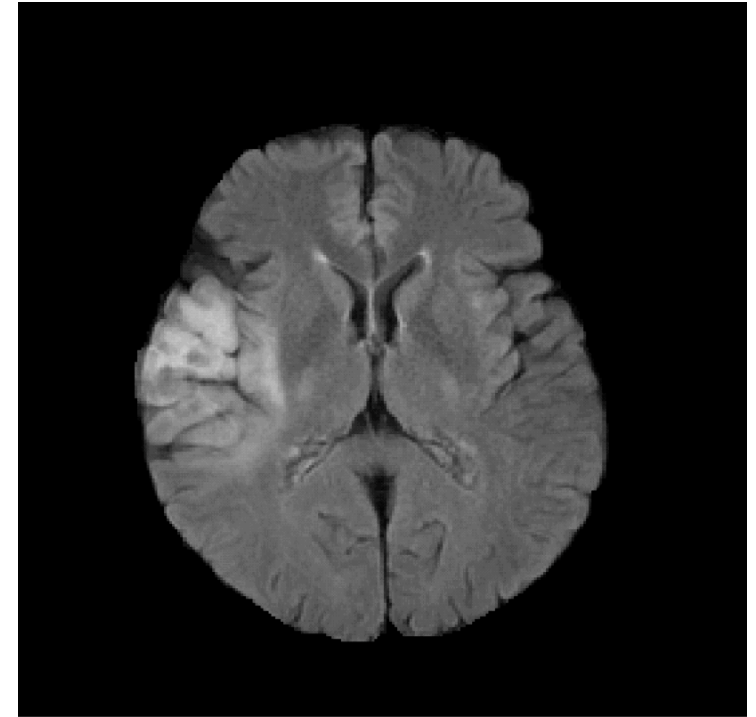
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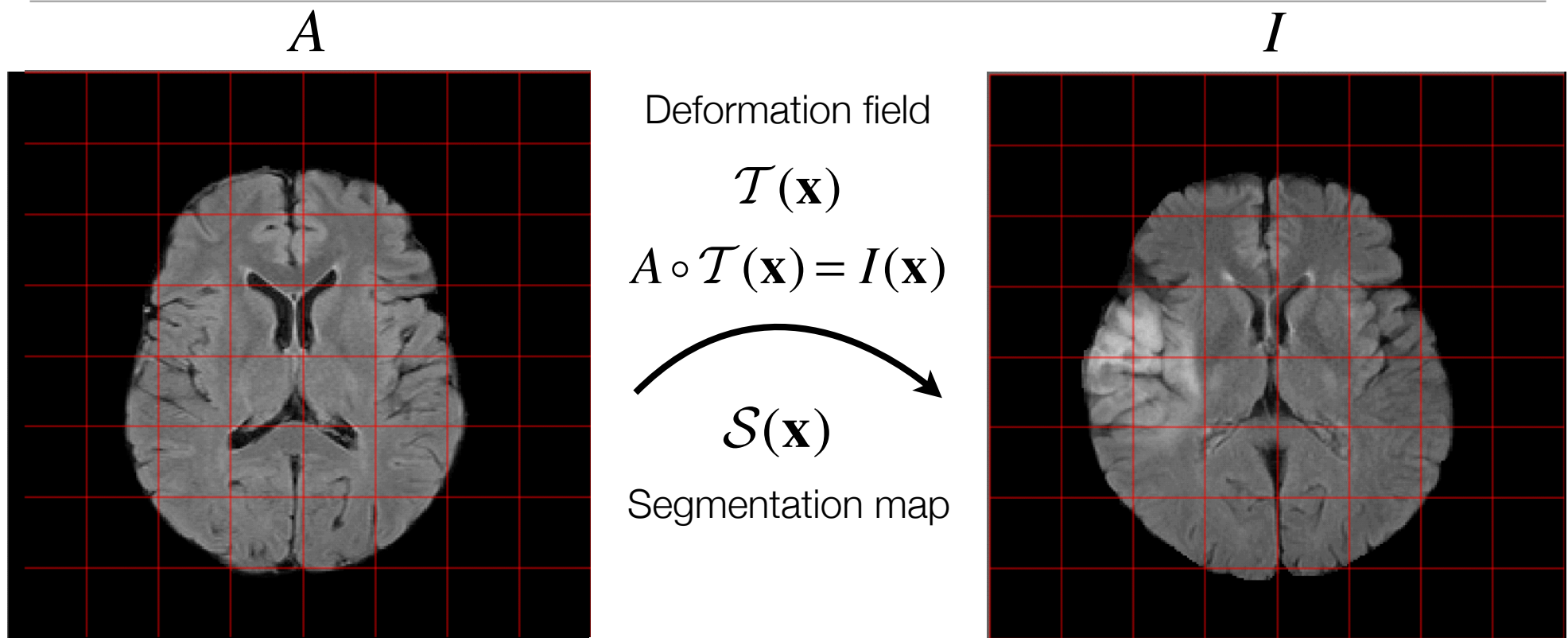
$$\mathcal{S}(\mathbf{x})$$

Segmentation map

$I$

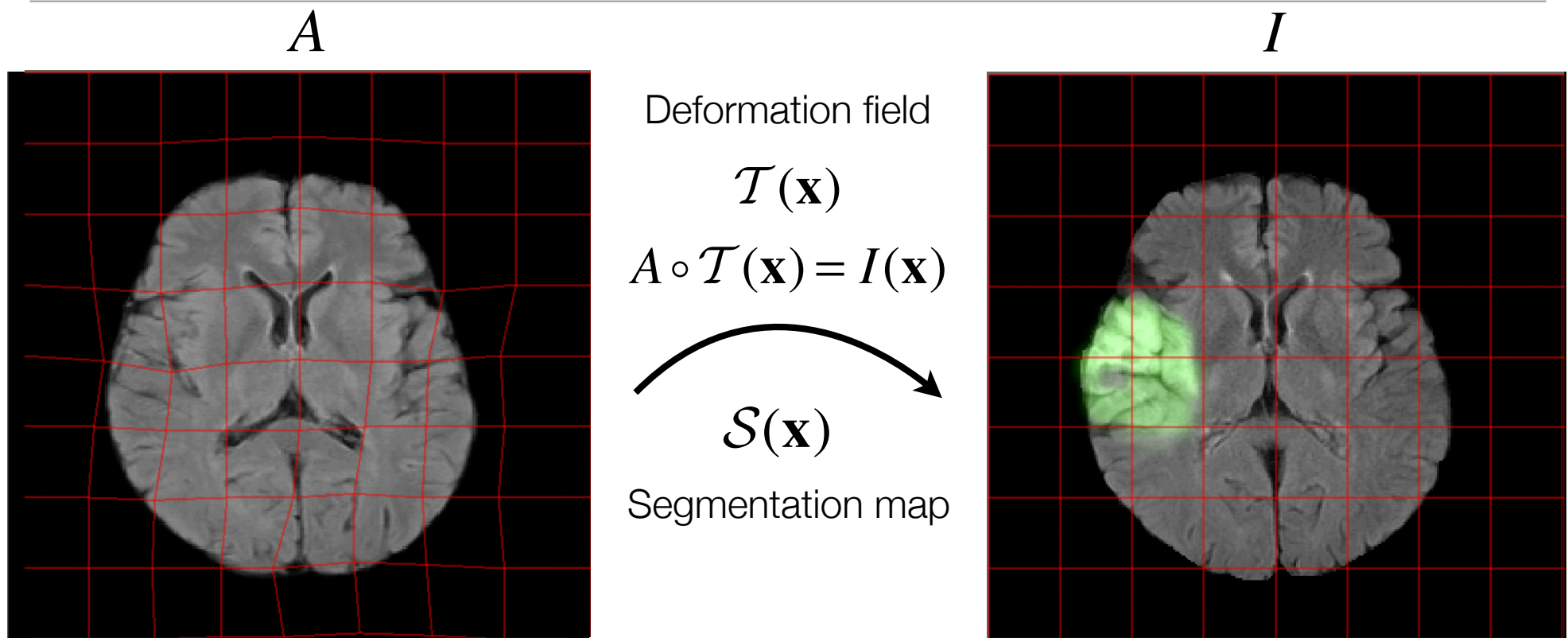


# Parametrization



- *Free form deformation* approach:
  - Deformation and Segmentation estimated on a grid  $\mathcal{G}$  superimposed to the images
  - Evaluation on the whole image by interpolation

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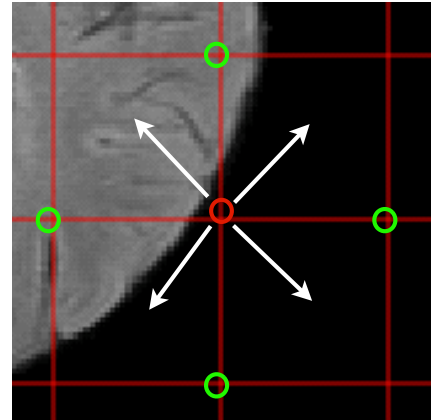
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# Markov Random Field Model

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- Assign to each grid node  $p$  a label  $l_p$  corresponding to a pair segmentation ( $s^l$ ) / displacement ( $\mathbf{d}^l$ )

$$l_p = \{s^{l_p}, \mathbf{d}^{l_p}\} \in \{0,1\} \times \{\mathbf{d}^1, \dots, \mathbf{d}^k\}$$

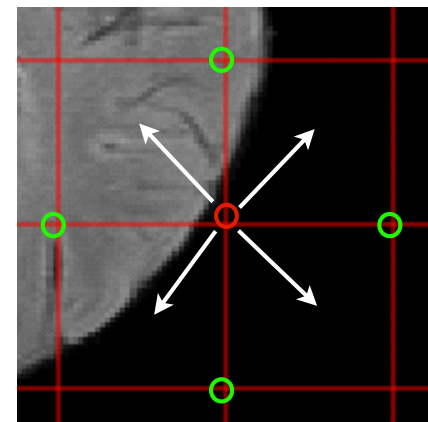


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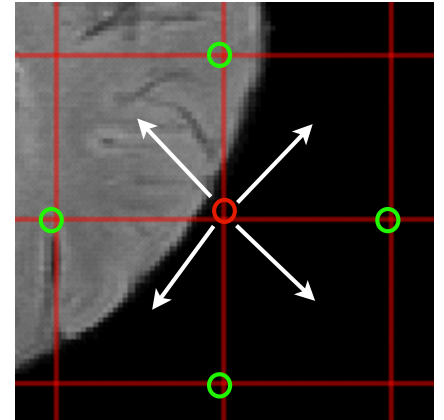
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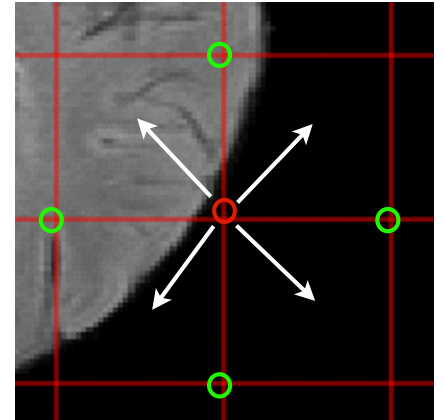
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- Markov Random Field Energy

$$E_{def,seg}(l) = \frac{1}{|\mathcal{G}|} \sum_{p \in \mathcal{G}} V_p(l_p) + \sum_{p \in \mathcal{G}} \sum_{q \in \mathcal{N}(p)} V_{pq}(l_p, l_q)$$



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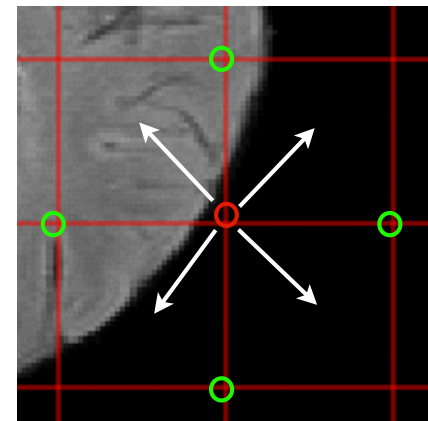
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Local consistency of the segmentation

Registration regularization



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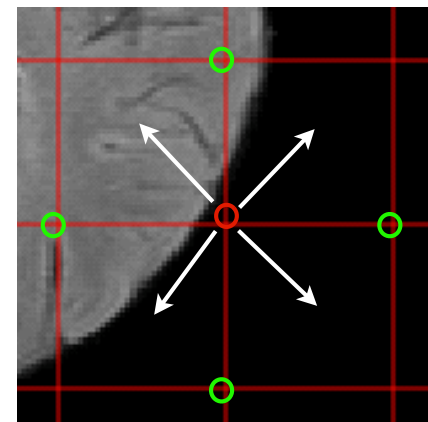
- Unary term

$$V_p(l_p) = \alpha V_{def}(l_p) + (1 - \alpha) V_{seg}(l_p)$$

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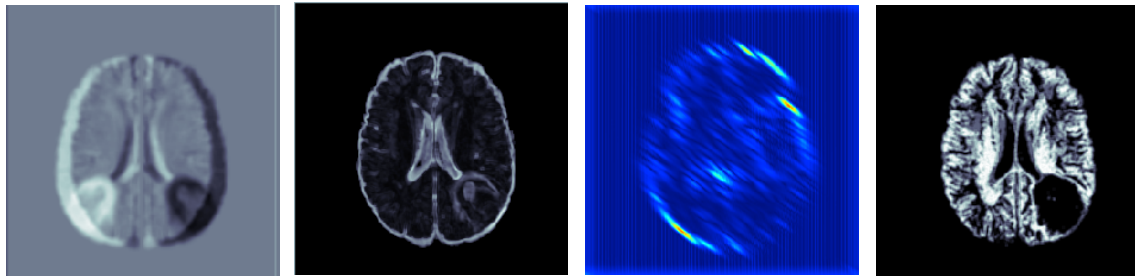
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# Unary Term: Segmentation

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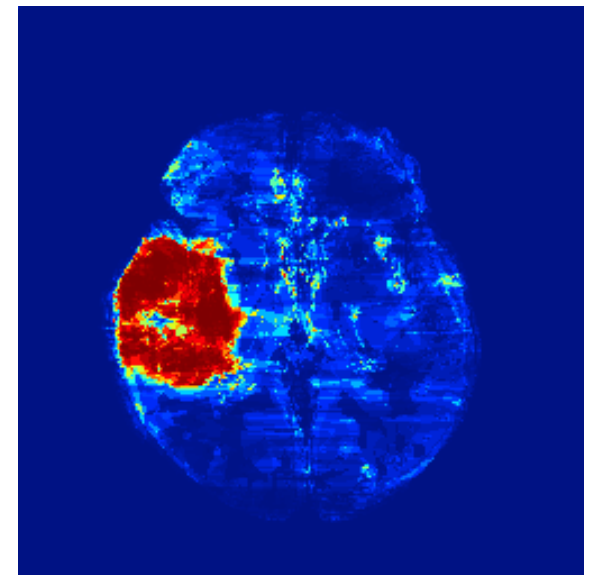
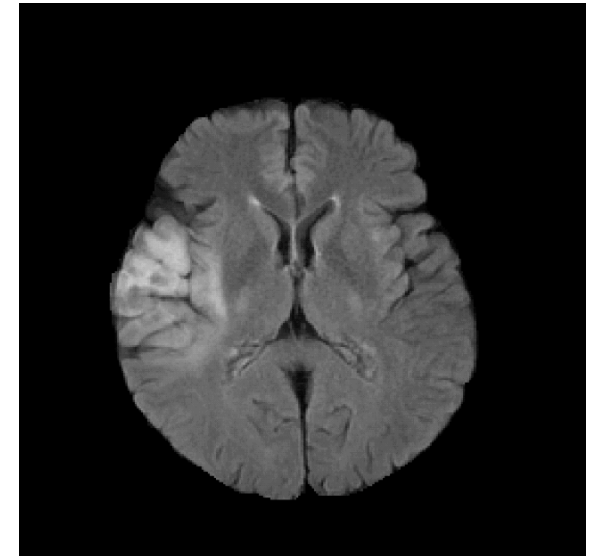
- Learning of a tumor vs background classifier
  - Features extracted from images



- Gentle Adaboost algorithm

Construction of a strong classifier as a combination of weak classifiers (decision stumps)

- **Any** classification technique can be used
- Nodes with **high classification probability** of being tumor should be labeled accordingly

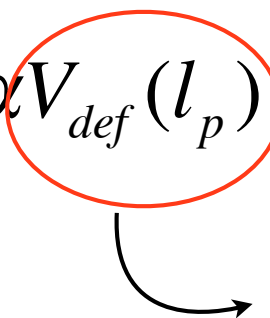


Boosting probability map

# Unary Term: Registration

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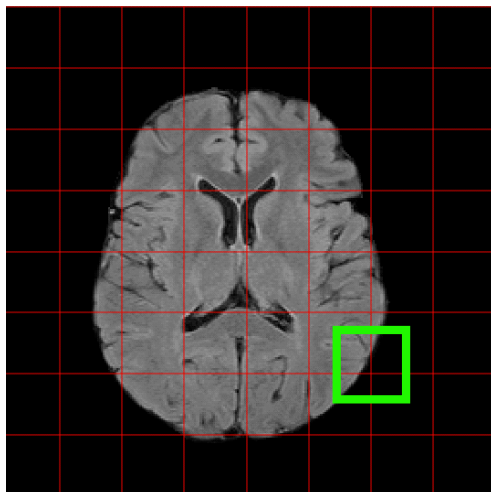
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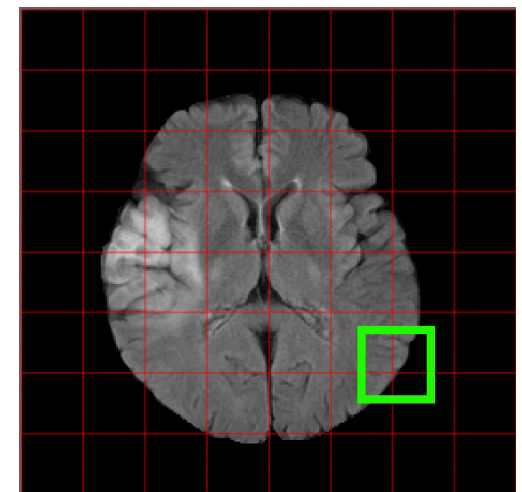
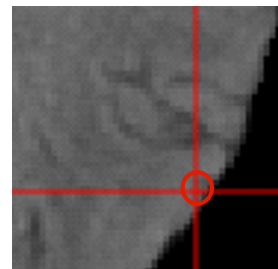
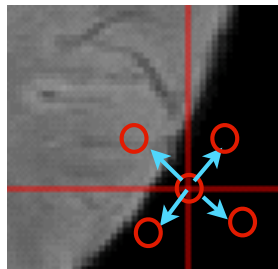
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- Outside the tumor area: Find correspondences

Compute the **similarity measure** for all possible displacements



*A*



*I*

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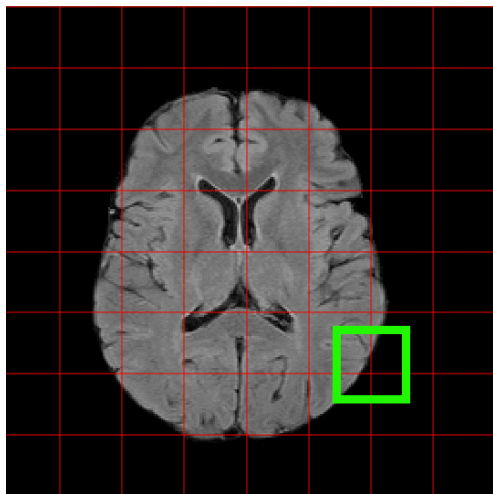
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$\swarrow$

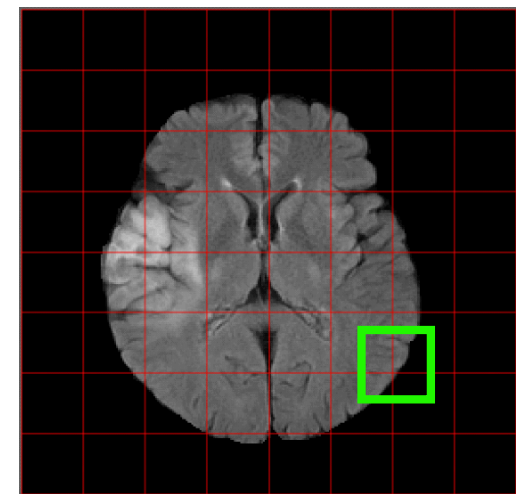
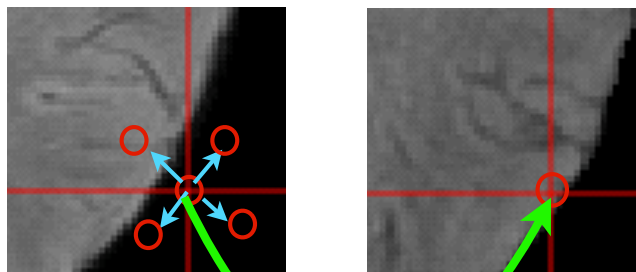
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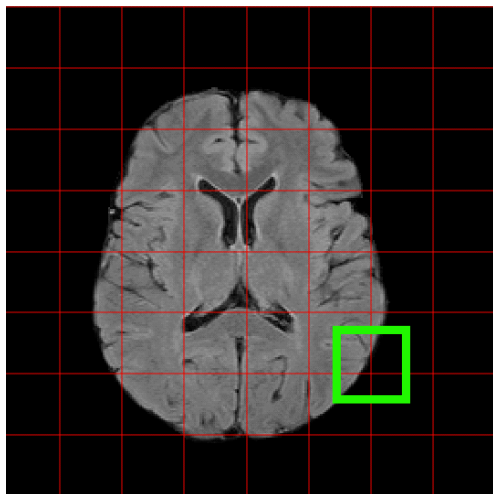
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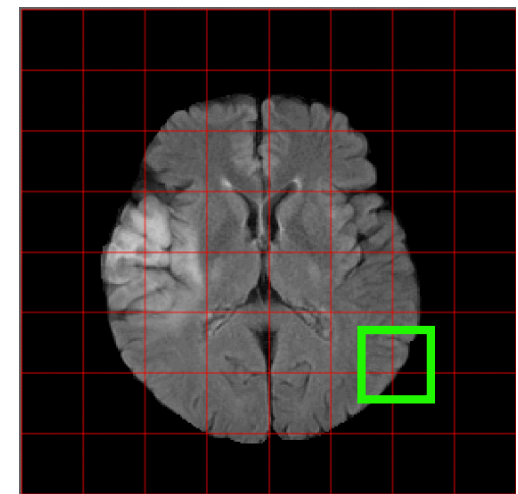
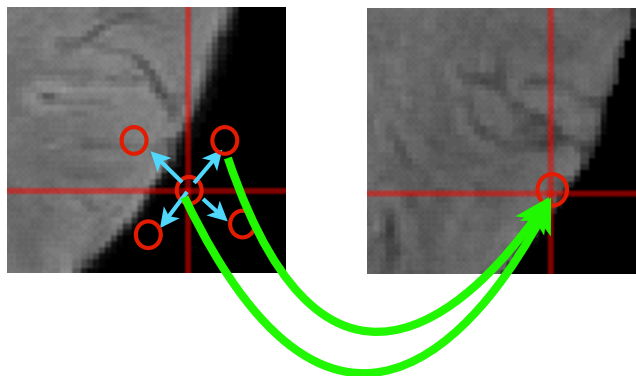
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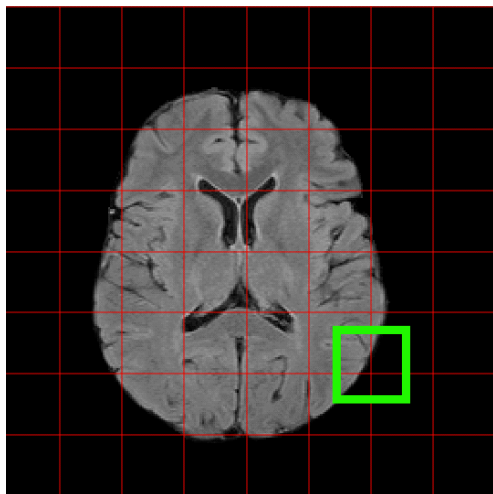
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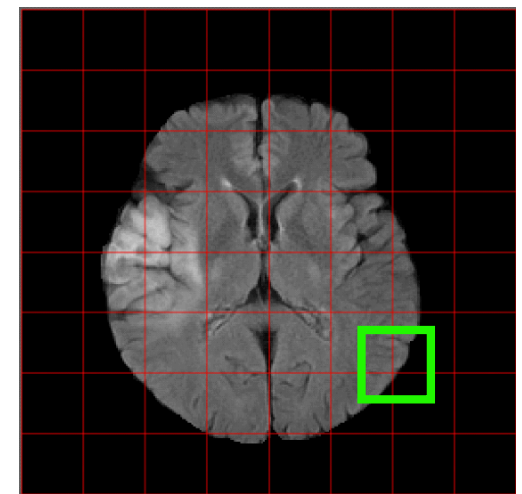
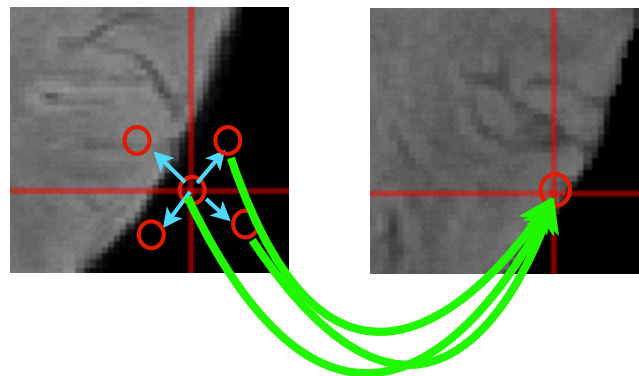
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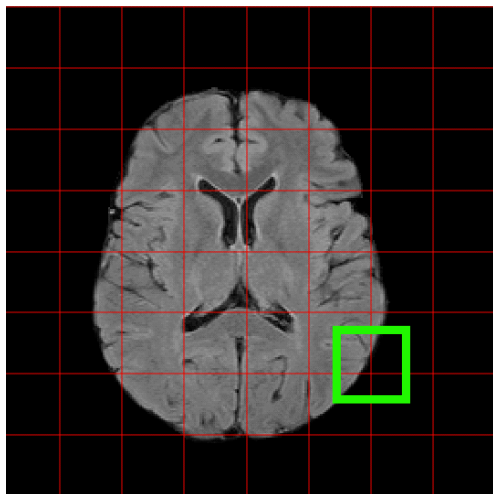
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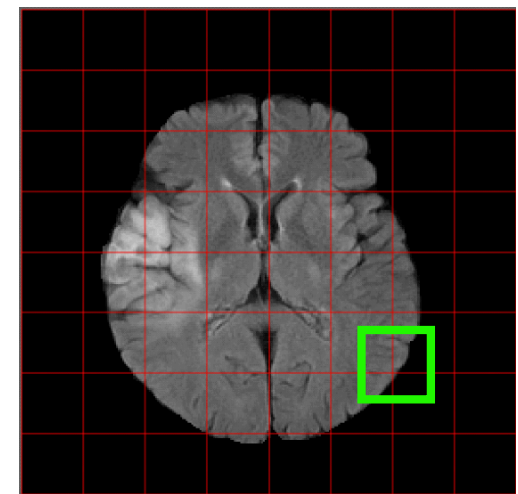
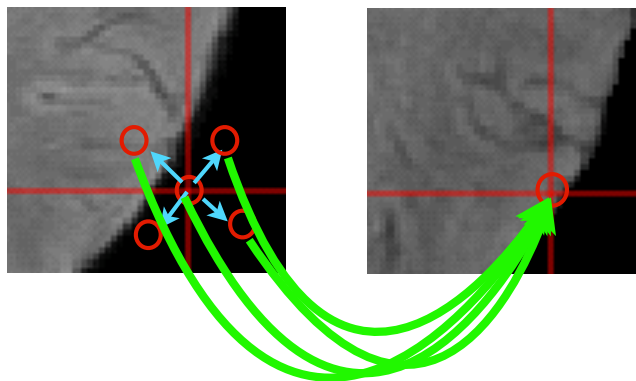
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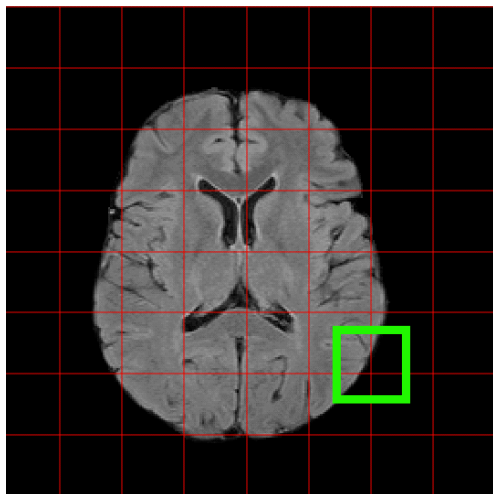
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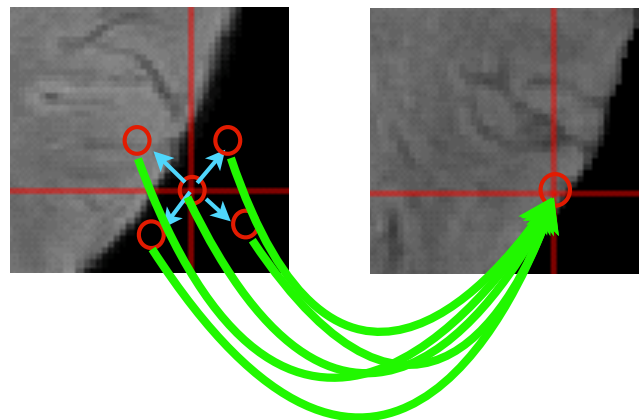
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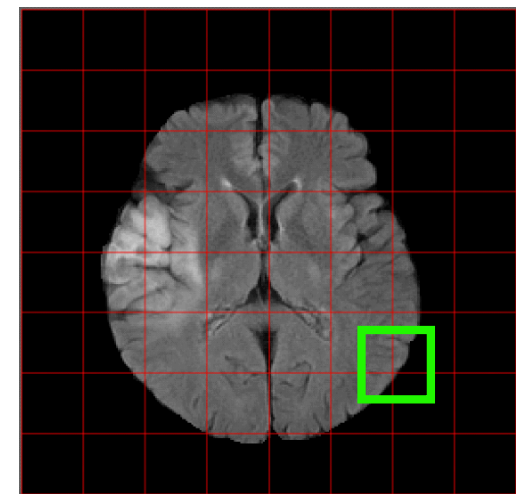
Compute the **similarity measure** for all possible displacements



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Similarity measure  $\text{Sim}(I(\mathbf{x}), A(\mathcal{T}(\mathbf{x})))$



$I$

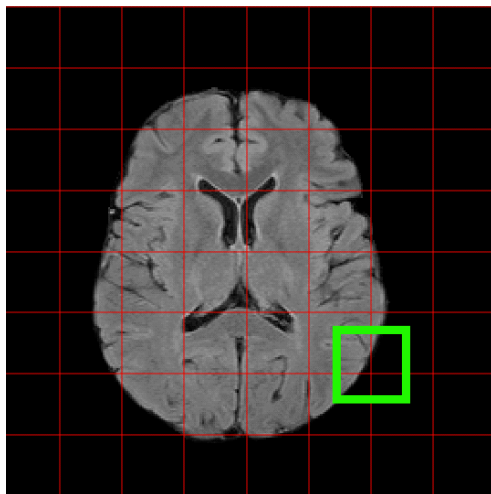
# Unary Term: Registration

$$V_p(l_p) = \alpha V_{def}(l_p) + (1 - \alpha) V_{seg}(l_p) \quad \text{Any kind of similarity criterion}$$

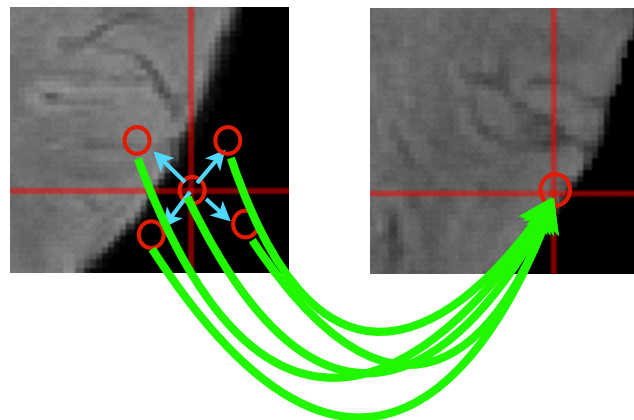
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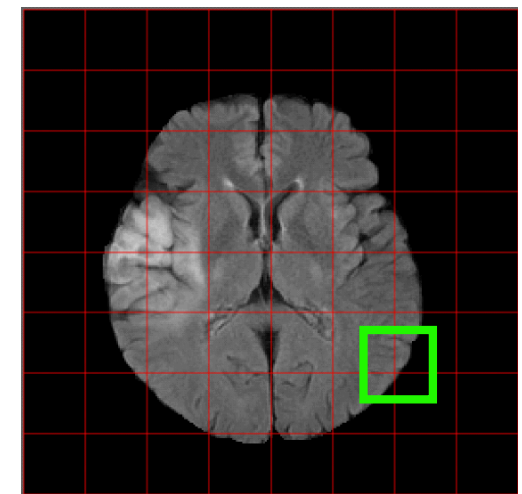
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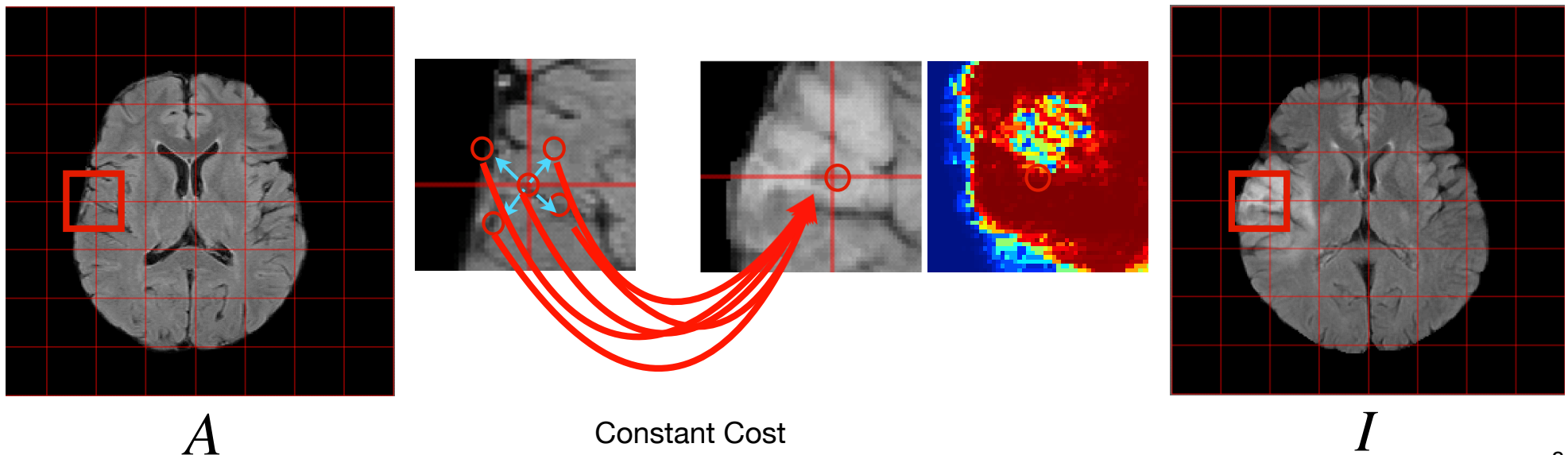
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- In the tumor area: No correspondences

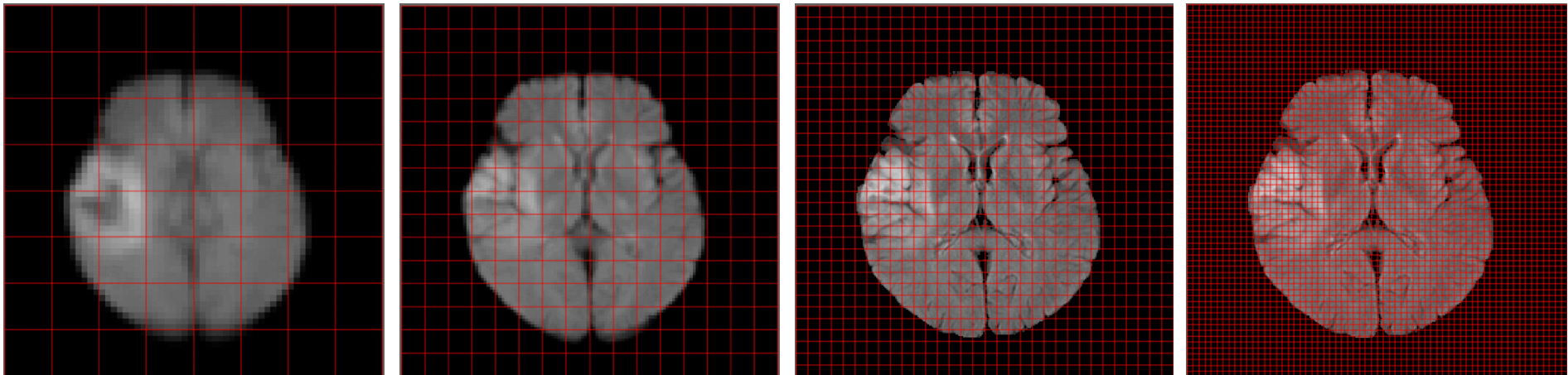
Constant cost independent of the displacement



# Implementation

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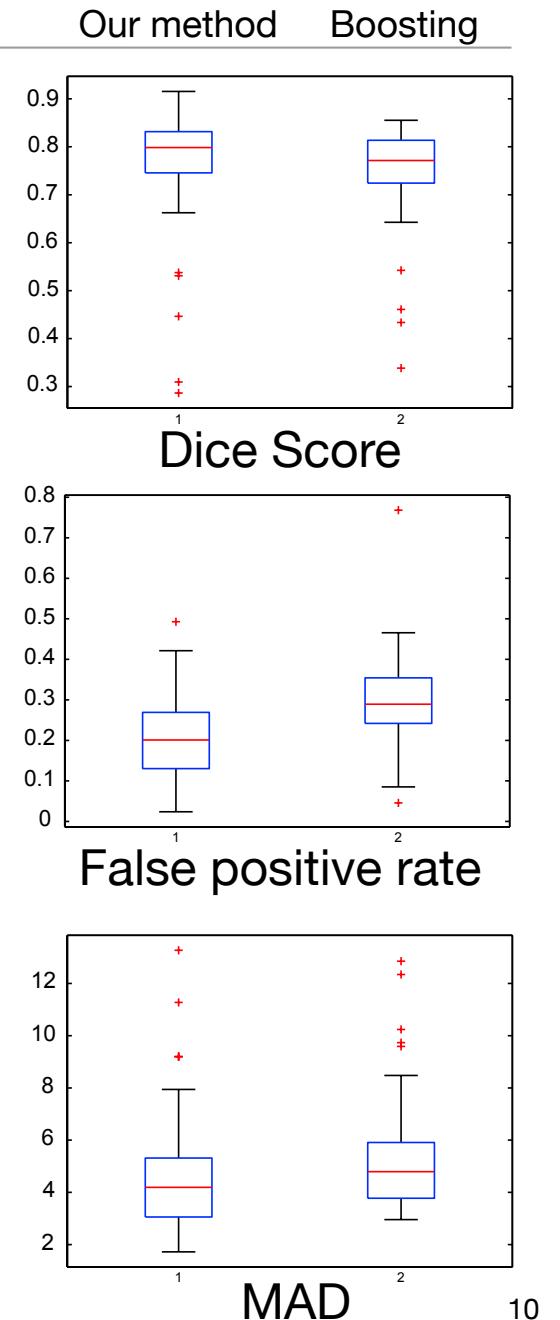
- Incremental Approach
  - 3 image levels, 4 grid resolutions



- Increasing influence of the segmentation (progressive diminution of  $\alpha$  value)
- Optimization
  - Linear programming (Komodakis et al. CVIU, 2008)
- Overall run time: 6 min (matlab implementation)

# Experimental Validation

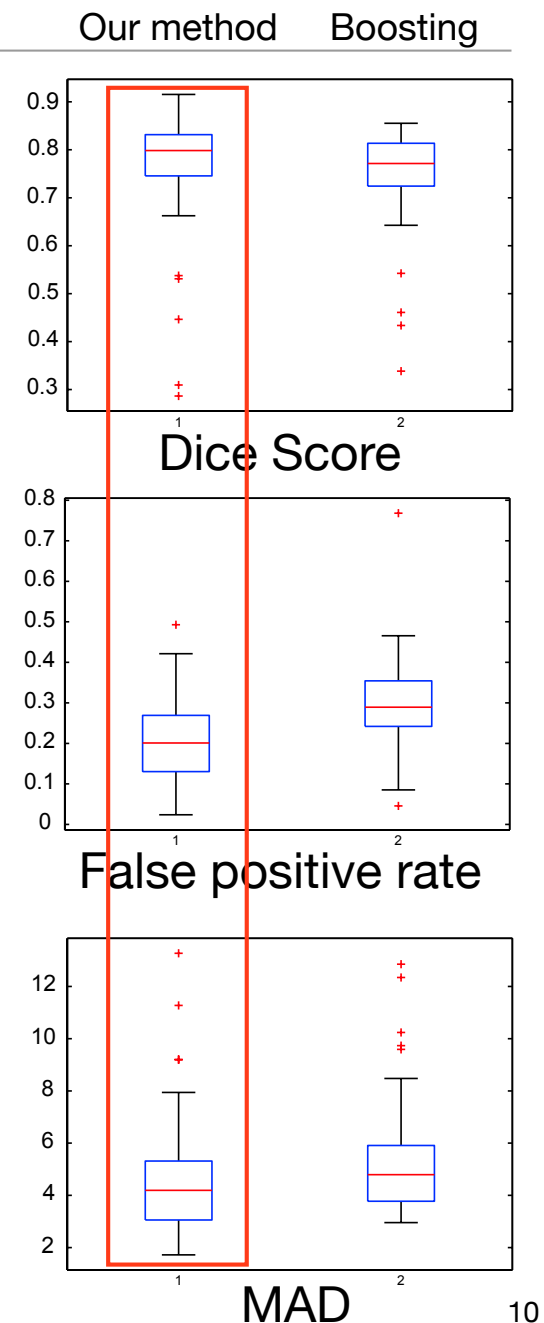
- Database: 97 T2 FLAIR volumes
  - Data likelihood learned on 40 volumes
  - Evaluation on 57 volumes
- Segmentation
  - Evaluated w.r.t manual segmentations
  - Compared with boosting classification with added pairwise smoothing (right on boxplots)
  - Median Dice: 77 to 80%, False positives: 30 to 20%, Mean Absolute Distance (MAD): 4.8 to 4.2mm
- Registration
  - Qualitative evaluation
  - Compared with Glocker et al. 2008, with masked pathology

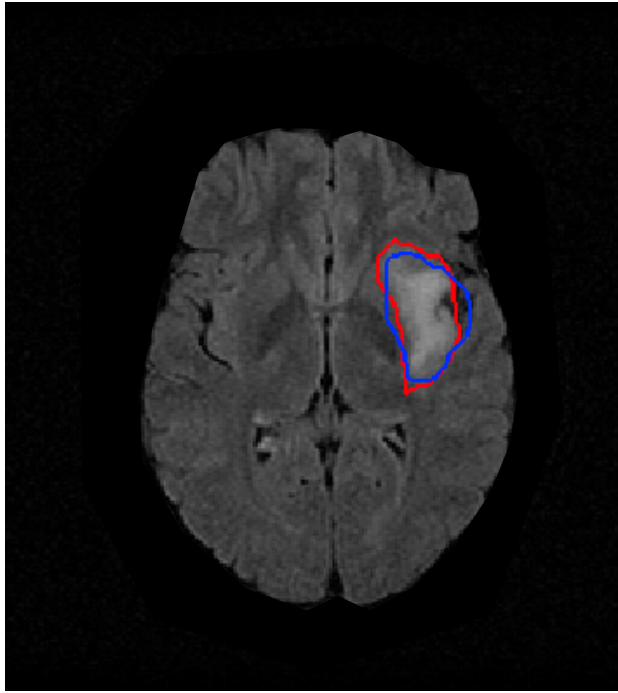
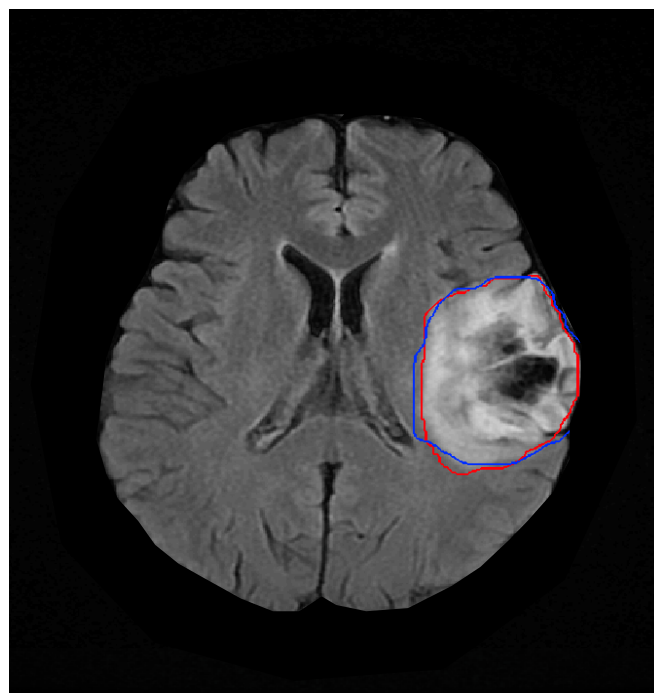
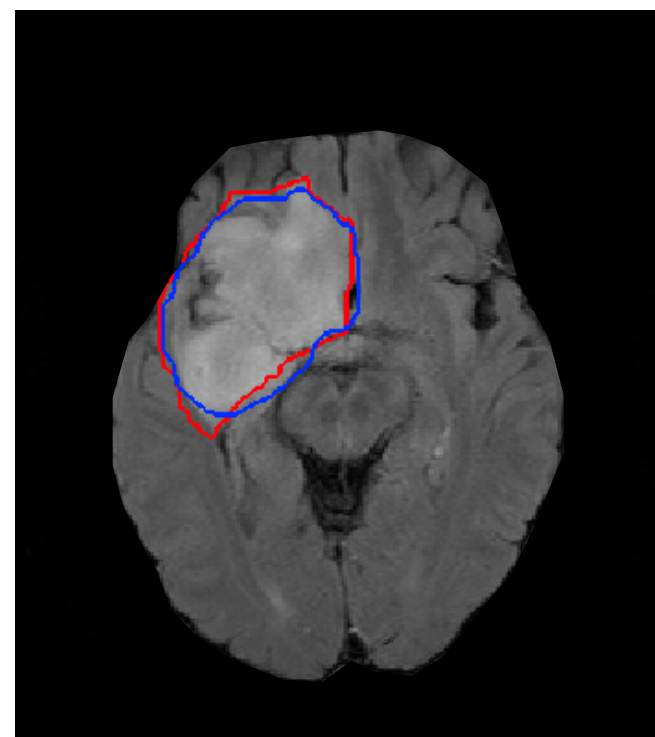
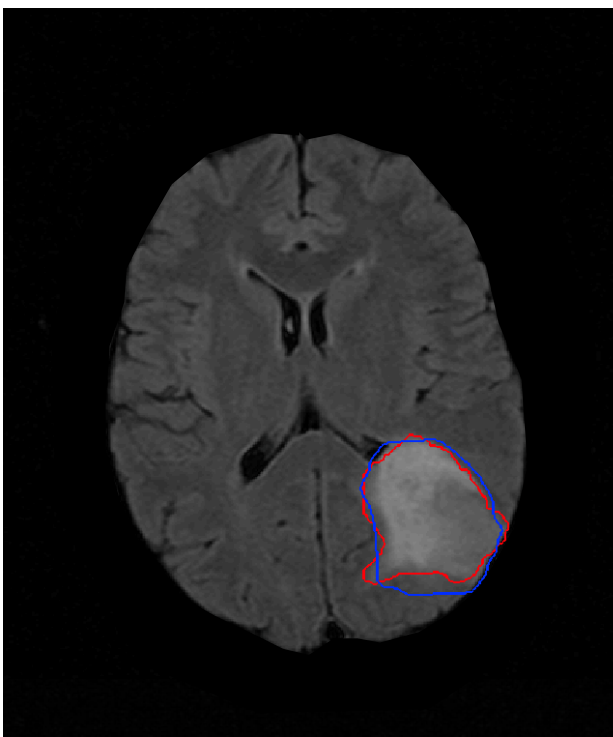
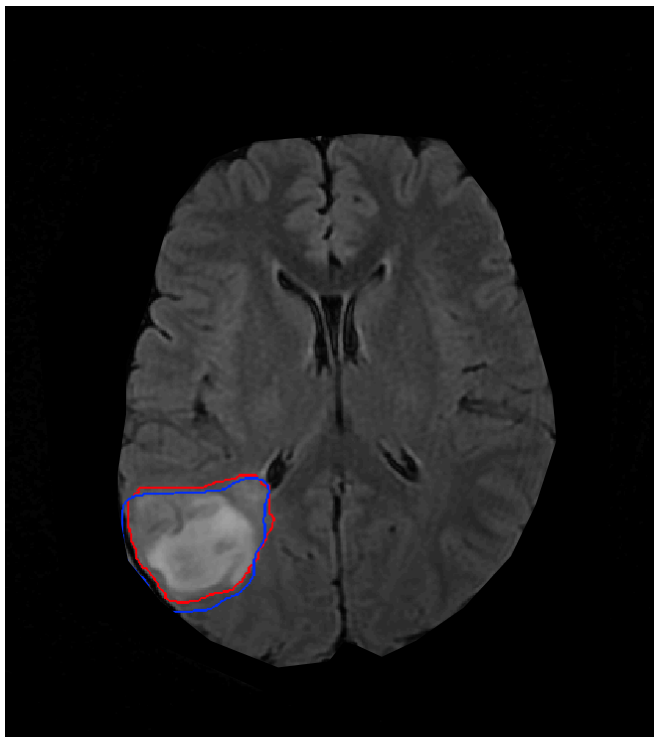




# Experimental Validation

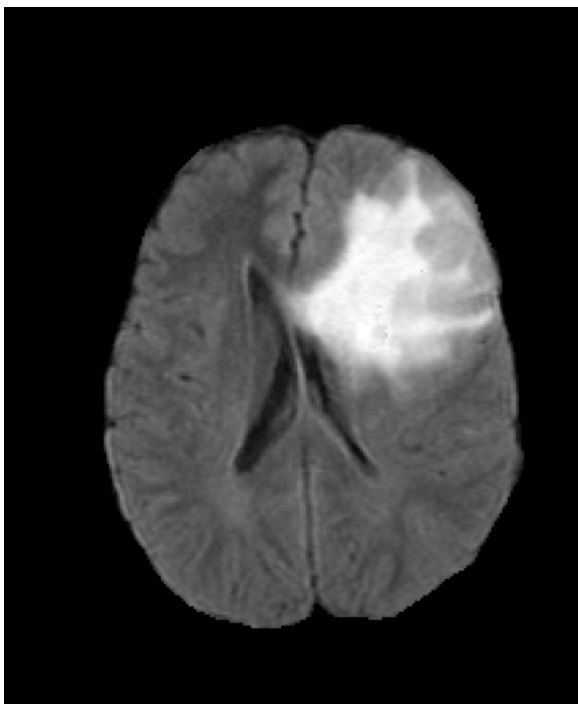
- Database: 97 T2 FLAIR volumes
  - Data likelihood learned on 40 volumes
  - Evaluation on 57 volumes
- Segmentation
  - Evaluated w.r.t manual segmentations
  - Compared with boosting classification with added pairwise smoothing (right on boxplots)
  - Median Dice: 77 to 80%, False positives: 30 to 20%, Mean Absolute Distance (MAD): 4.8 to 4.2mm
- Registration
  - Qualitative evaluation
  - Compared with Glocker et al. 2008, with masked pathology



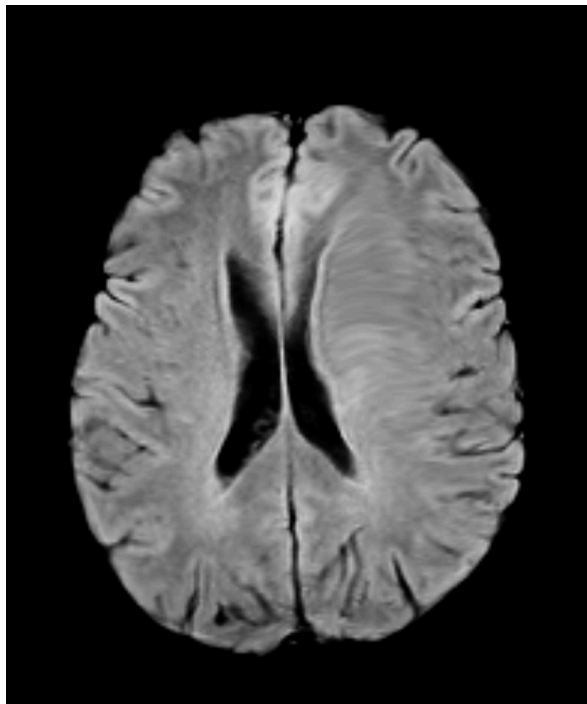


Red: Ground truth

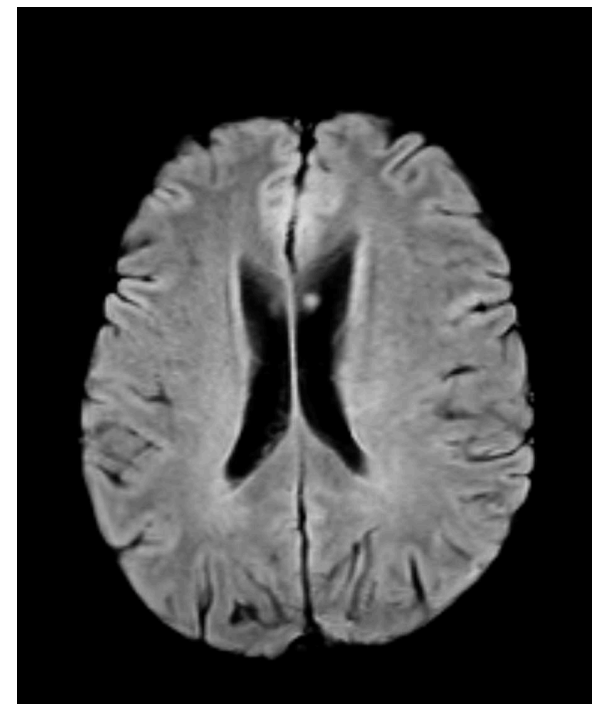
Blue: Automatic segmentation



Original image



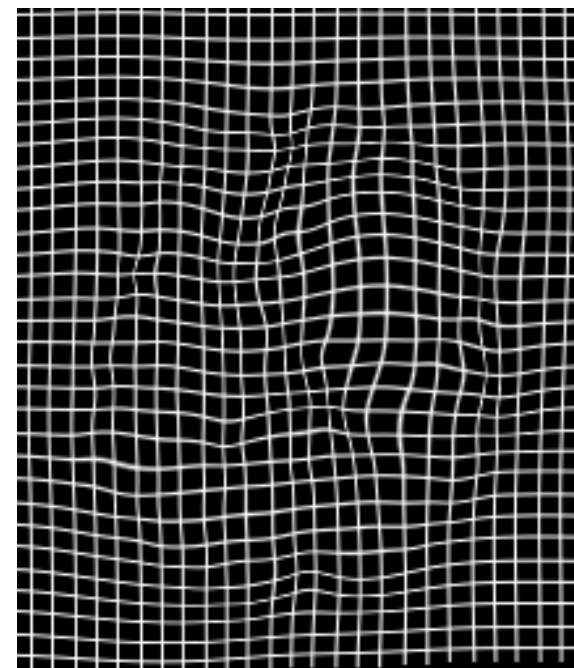
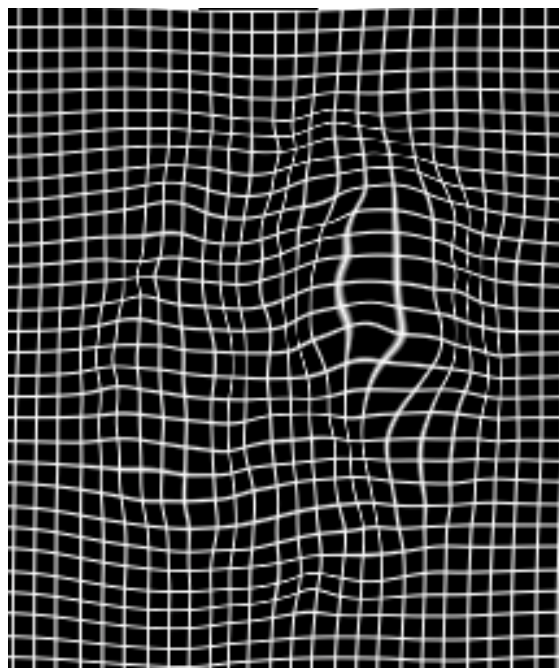
Glocker et al. 2008  
Deformed image



Our method  
Deformed image

Left:  
Glocker 08  
Deformation field

Right  
Our method  
Deformation field



# Conclusion

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- Simultaneous registration and segmentation method
- Modular w.r.t image modality, similarity criterion and classification technique
- Can be adapted to any clinical context
- Fast and efficient optimization (ongoing work to reduce the run time to a few seconds)
- State of the art results
- Future work
  - Local spatial position prior information
  - Registration uncertainties
  - Adaptation to registration/segmentation before and during surgery with tumor resection

Poster *Th-1-AG-14*

Thursday 13:30-15:00

Thank you for your attention

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Questions?