

## ABSTRACT

We present an efficient, unsupervised FLAIR-only MS lesion segmentation algorithm that does not use tissue priors or parametric models. An edge-based model of partial volume averaging (PVA) is used to estimate fuzzy membership profiles in order to map FLAIR graylevel to tissue class.

## INTRODUCTION

### Automated Segmentation of MS Lesions:

- Can improve speed, accuracy, and reproducibility
- Facilitates large, retrospective, and temporal studies

### Approaches and Challenges to Automation:

- Parametric models & supervised classification:
  - Challenged by variability in anatomy, and lesion load, as well as MRI scanner and sequence parameters
  - Partial volume averaging (PVA) not adequately modeled
- If require multiple MRI modalities:
  - Need accurate registration
  - Additional scan time and cost
  - Limited retrospective image availability

## EDGE-BASED PVA MODEL

### Inspiration

- Image edges correspond to PVA-affected voxels: Figure 1.
- Edge is proportional to change in partial volume fraction ( $\alpha$ )

### Model of PVA Graylevels:

$$y_{ik}(x) = \alpha_{jk}(x) \times y_j(x) + (1 - \alpha_{jk}(x)) \times y_k(x) \quad \begin{cases} y_j(x) \sim p_j(y) \\ y_k(x) \sim p_k(y) \\ \alpha_{jk}(x) \in [0,1] \end{cases}$$

### Estimation of Partial Volume Fraction (PVF)

We desire global estimate of PVF given graylevel:  $\alpha(y)$

- Estimate change in PVF in  $x$  domain using edge image:

$$\alpha'(x) \Leftarrow |\nabla_x(Y(x))|$$

- Migrate to the global  $y$  domain using the conditional expectation, with KS density estimator:

$$\alpha'(y) = p(\alpha'(x) = 1 | y)$$

- Identify pure tissue classes  $y_j, y_k, \dots$  as minima in  $\alpha'(y)$ :

Figure 2: top

- Integrate to between  $y_j$  and  $y_k$  to resolve  $\alpha(y)$ :

$$\alpha_{jk}(x) = \frac{\int_{y_k}^y \alpha'(y) dy}{\int_{y_k}^{y_j} \alpha'(y) dy}, \quad y \in [y_k, y_j]$$

Figure 2: middle

- Construct fuzzy membership profiles  $\xi(y)$ :

$$\xi(c_k | y) = \begin{cases} 0, & y \in [0, y_j) \\ \alpha_{jk}(y), & y \in [y_j, y_k) \\ 1, & y = y_k \\ 1 - \alpha_{jk}(y), & y \in (y_k, y_\ell] \\ 0, & y \in (y_\ell, y_{max}] \end{cases}$$

Figure 2: bottom

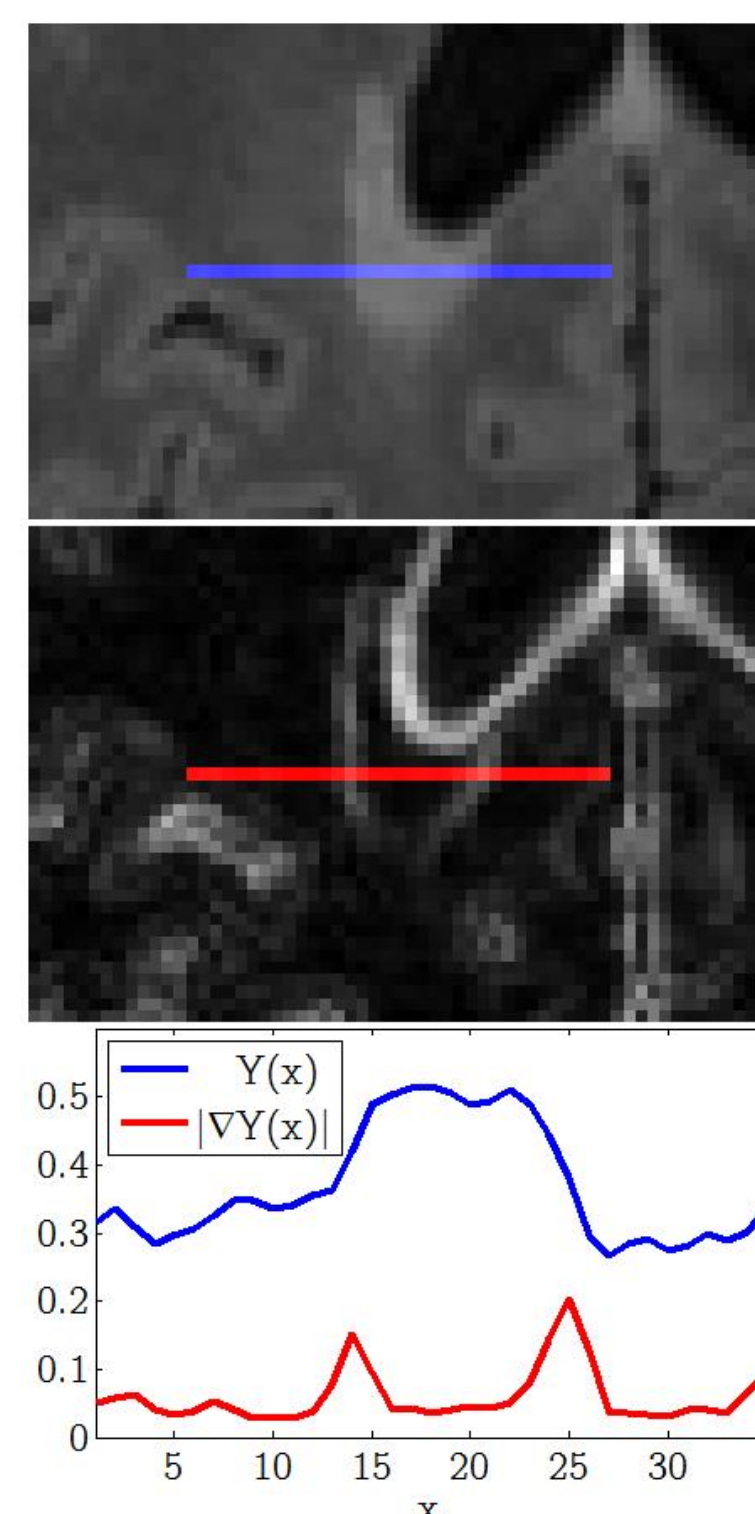


Figure 1. Example 1D profiles through FLAIR image  $Y(x)$  and edge image  $E(x)$  showing correlation of edge with change in PVF.

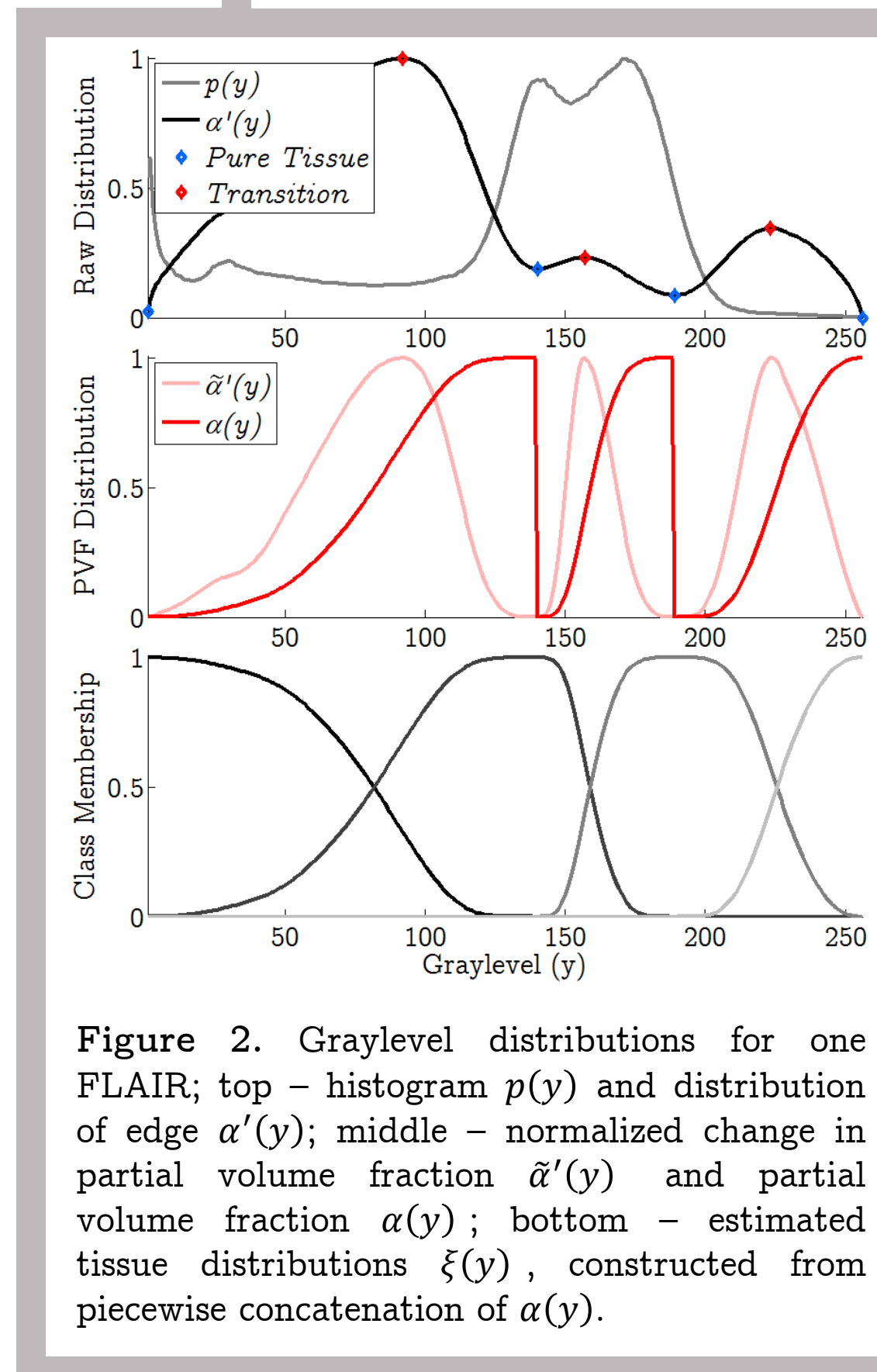
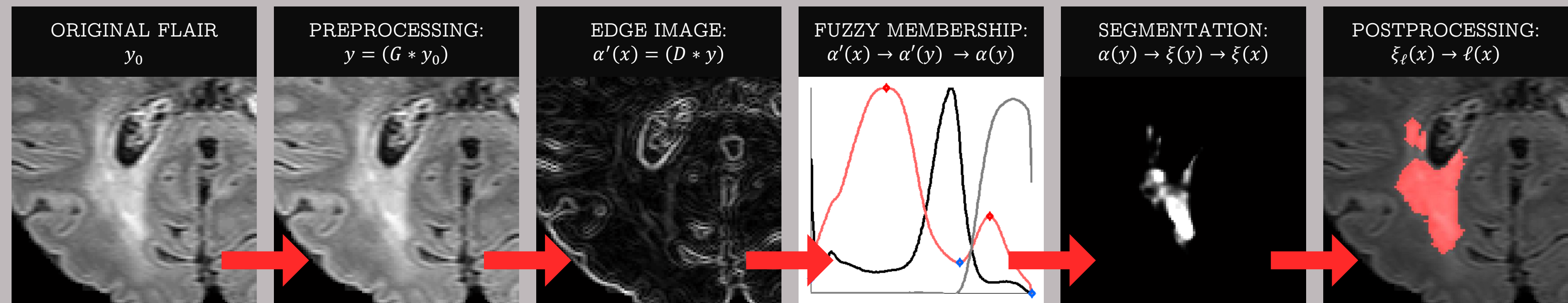


Figure 2. Graylevel distributions for one FLAIR; top - histogram  $p(y)$  and distribution of edge  $\alpha'(y)$ ; middle - normalized change in partial volume fraction  $\hat{\alpha}'(y)$  and partial volume fraction  $\alpha(y)$ ; bottom - estimated tissue distributions  $\xi(y)$ , constructed from piecewise concatenation of  $\alpha(y)$ .



## MODEL CHARACTERISTICS

### Advantages:

- Unimodal: FLAIR only
- Nonparametric: no assumed distributions
- Registration-free
- Sub-voxel precision
- Unsupervised

### Assumptions / Requirements:

- Univariate separability in FLAIR \*
- Bias corrected \*
- Brain extracted
- Homogeneous lesion appearance

\* future works to overcome these

## RESULTS

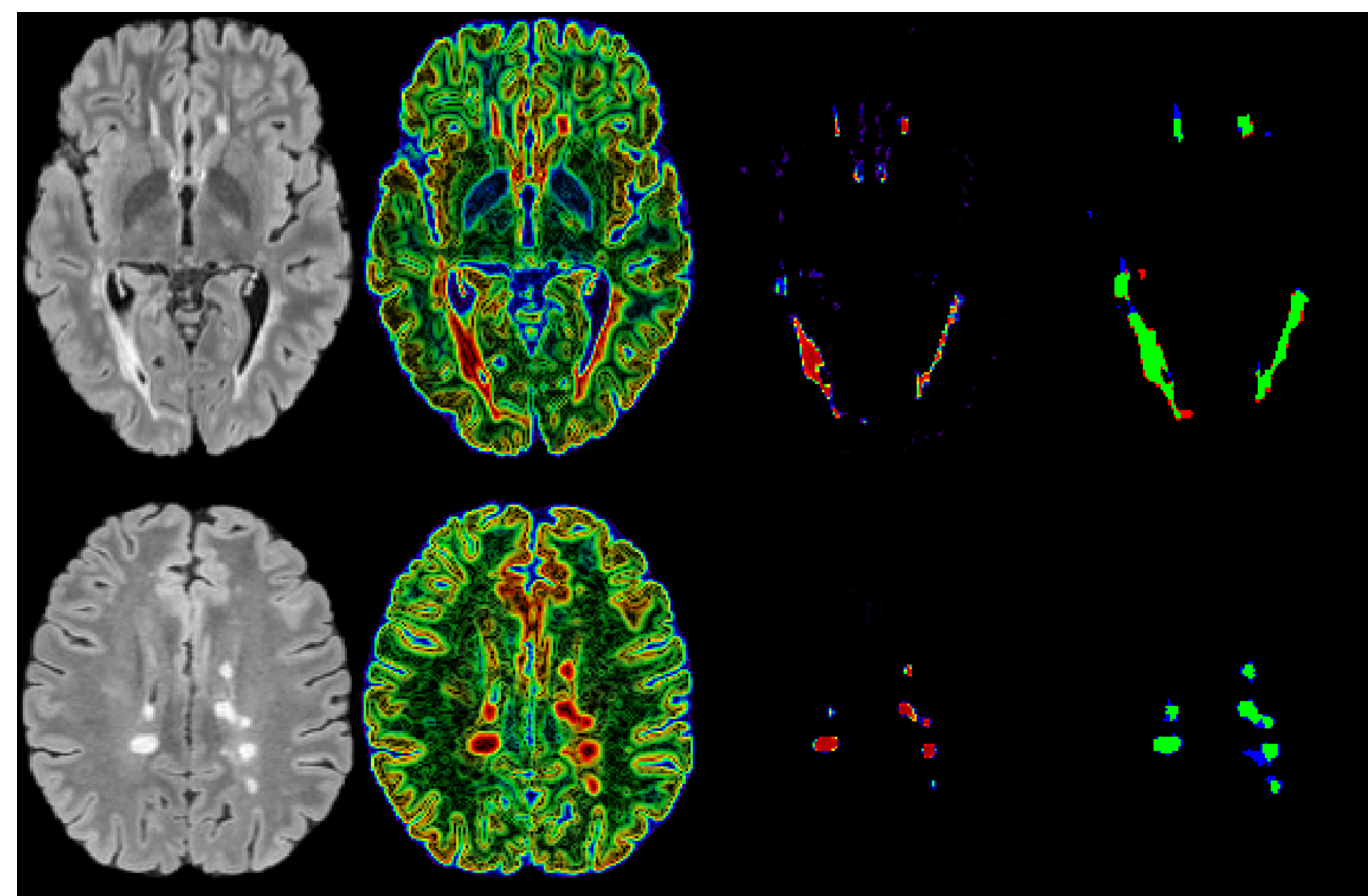


Figure 3. Example edge images and segmentations: (a) original FLAIR; (b) edge image with brightness depicting edge magnitude and pseudocolour depicting FLAIR graylevel (note orange PVA voxels); (c) pseudocolour initial fuzzy segmentation; (d) segmentation results, green is TP, red is FP, blue is FN.

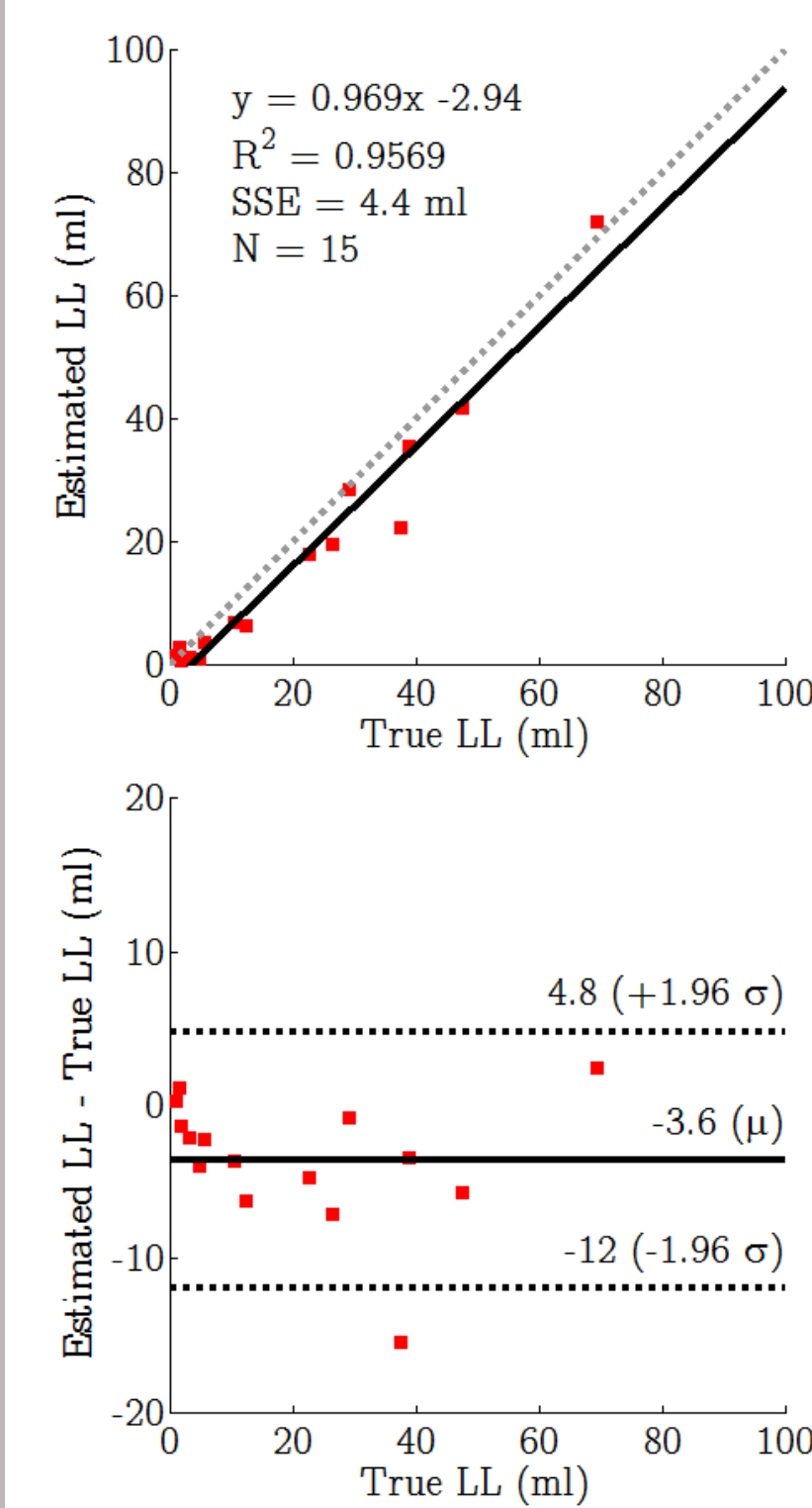


Figure 4. Bland-Altman plots showing good lesion volume agreement with expert segmentations, with a slight bias towards undersegmentation.

### Segmentation Performance

- Mean DSC = 0.60 (all volumes)  
= 0.70 (lesion load > 5 ml)
- Mean PPV = 0.80; TPR = 0.53

### Volume Agreement

- Pearson's  $R^2 = 0.96$
- Undersegmentation bias (slope = 0.97)

### Processing Speed

30 seconds per volume  
(MATLAB R2016, Intel i7, 16GB RAM)

### ANCOVA:

- DSC correlated with lesion load ( $p = 0.013$ )
- DSC uncorrelated with scanner ( $p = 0.354$ )

## FUTURE WORKS

### Local Analysis:

- Compute  $\alpha'(y)$  locally (for each voxel), then estimate pure tissues graylevels  $\bar{y}_c(x)$
- Estimate bias field from  $\bar{y}_c(x)$  by fitting low order basis functions
- Overcome: (1) bias field, (2) overlapping graylevel distributions (bright GM)

### Defining “Pure Lesion”

- Small, inhomogeneous lesions have less prominent extrema in  $\alpha'(y)$
- Multi-scale edge analysis could possibly resolve this

### Prior-Free Ventricle Segmentation

- Flow-through artifacts could additionally be excluded

## REFERENCES

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## AKNOWLEDGEMENTS



## PRE- & POST-PROCESSING

### Preprocessing

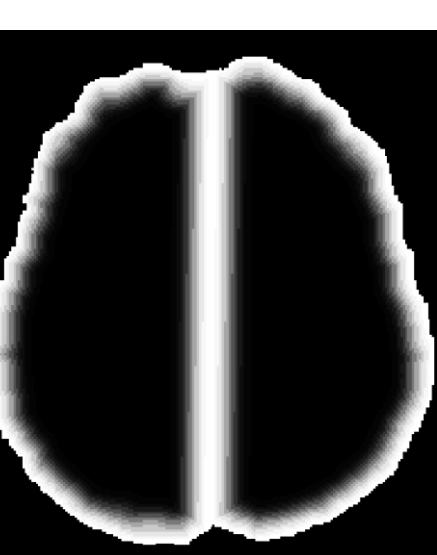
- Bias correction using SPM12: “Segment”
- Brain masks: provided by MSSEG Competition

### Region Growing

- Mimic radiologist: liberal lesion borders
- Compare candidate voxel at  $x^*$  to  $q^{th}$  quantile of  $y$  in initial segmentation  $A$ :  
if:  $|y(x^*) - Q_{y \in S}(q)| < \psi$  then:  $A \leftarrow A \cup x^*$

### Parameter Optimization

- Objective: mean DSC
- Parameters:
  - Fuzzy threshold  $\tau$
  - FPR distances & minimum lesion size
  - Region growing parameters



### Postprocessing

- Threshold fuzzy segmentation:  $\xi(x) \geq \tau$
- Enforce minimum lesion size
- Distance from brain edge and midline (above left)