

Effectiveness of Intensity Normalization on MRIs of Human Brains with Multiple Sclerosis

Mohak Shah^{1,3}, Yiming Xiao¹, Simon Francis^{2,3}, Douglas Arnold^{2,3}, D. Louis Collins²,
Tal Arbel¹

¹Centre for Intelligent Machines, McGill University

²Montreal Neurological Institute, McGill University

³NeuroRx Research, Montreal

Workshop on Medical Image Analysis on Multiple Sclerosis
MIAMS-2009

September 20, 2009

Mohak Shah

Outline

- ☐ Introduction
- ☐ Nyul Intensity Normalization
- ☐ Evaluation of Nyul approach
- ☐ Qualitative and Quantitative evaluation
- ☐ Conclusion and Future work

The Need for Intensity Normalization

- Variations in MRIs
 - Acquisition protocols;
 - Multi-site scans; scanner manufacturers; models from same manufacturer;
 - Effect of pathology; Varying stages of disease
- Lack of standard scale makes the generalization of intensity behavior difficult
- Reliance on assumptions over distribution of tissue intensities
 - large variations in intensity ranges can violate these

The Need for Intensity Normalization

- These variations are accounted for by
 - Standard acquisition protocols
 - Intensity in-homogeneity correction
 - Intensity Normalization (Standardization)

The Need for Intensity Normalization

- These variations are accounted for by
 - Standard acquisition protocols
 - Intensity in-homogeneity correction
 - **Intensity Normalization (Standardization)**

Approaches to Intensity Normalizations

☐ Various Approaches

- (Christensen, 1996); (Weisenfeld and Warfield, 2004); (Hellier, 2003); (Jäger *et al*, 2006)

☐ Main drawback

- Accuracy vs. Speed vs. Adaptability

☐ Nyul Approach (Nyul et. al., 1999, 2000)

- Best of both worlds
- One of the most widely approaches

Nyul Intensity Normalization (Nyul *et al.* 1999, 2000)

☐ Extensive Evaluation

- available on inter- and intra-patient variations
- But not on multi-site multi-scanner multi-spectral data
- Neither in the presence of pathology

Nyul Intensity Normalization (Nyul *et al.* 1999, 2000)

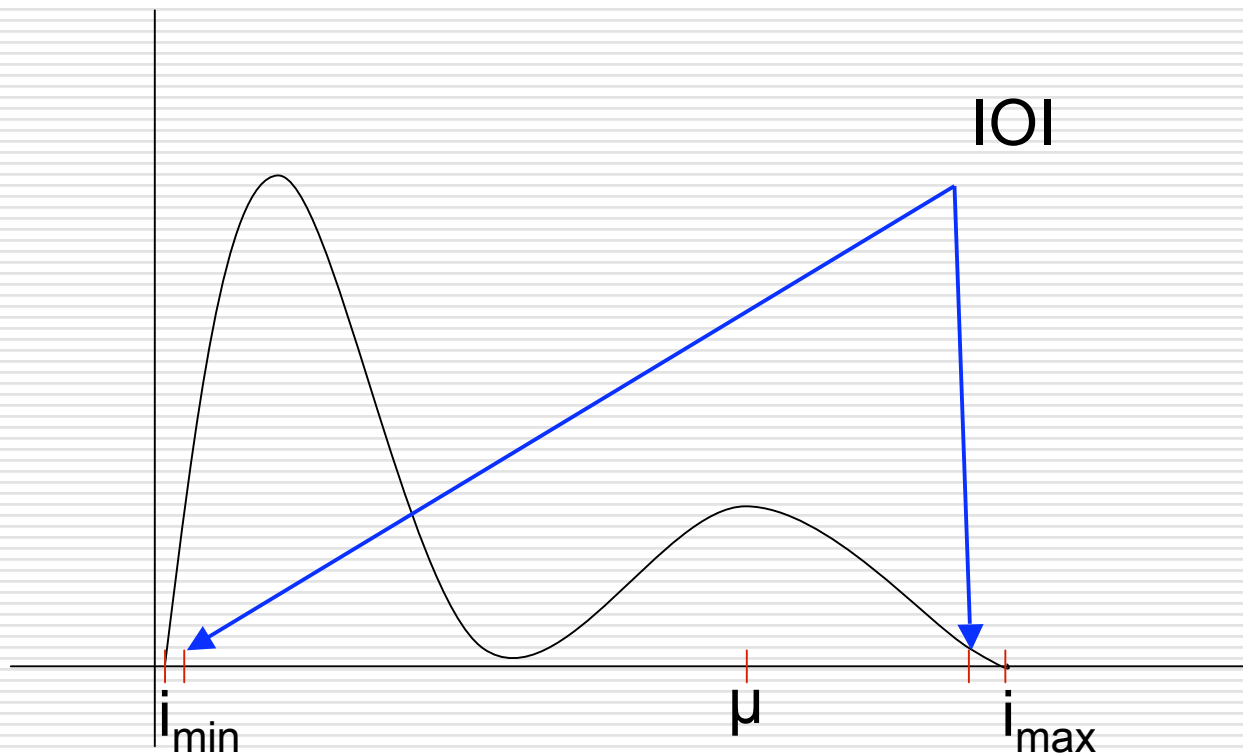
□ Extensive Evaluation

- available on inter- and intra-patient variations
- But not on multi-site multi-scanner multi-spectral data
- Neither in the presence of pathology

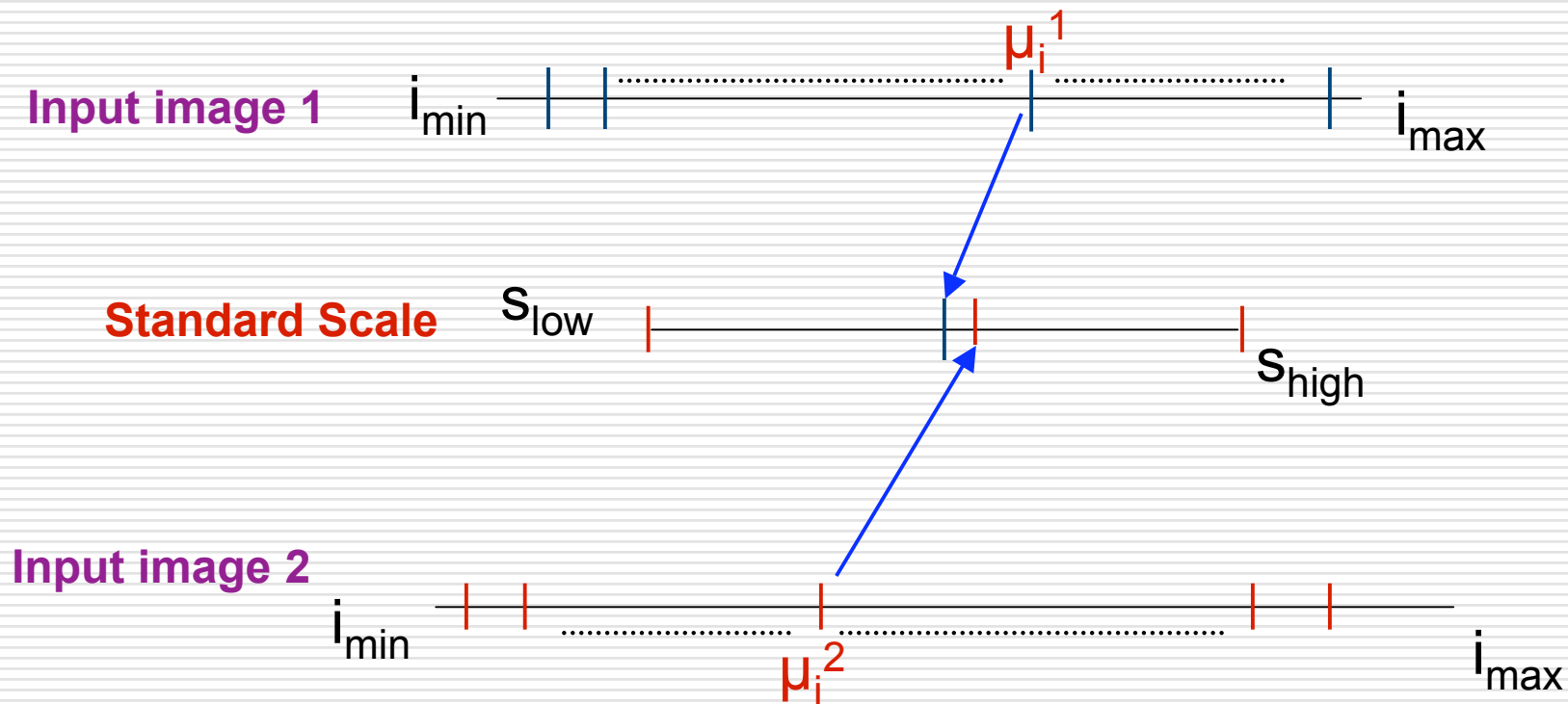
Nyul Approach in a nutshell

- ❑ Decile Formulation (Nyul *et al.* 2000)
- ❑ Two stage approach
- ❑ Training
 - Input a value range for forming a standard scale
 - Determine standard scale landmarks (mapping points) from the training data
- ❑ Transformation
 - Use standard scale for a piece-wise linear mapping of input image to standard scale

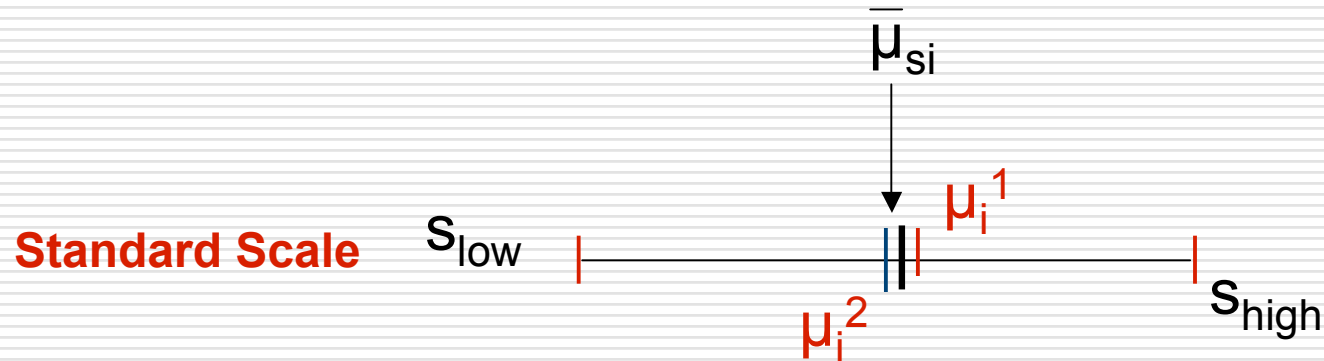
The Intensity Normalization (Training): Intensity of Interest (IOI)



The Intensity Normalization (Training): mapping to Standard scale

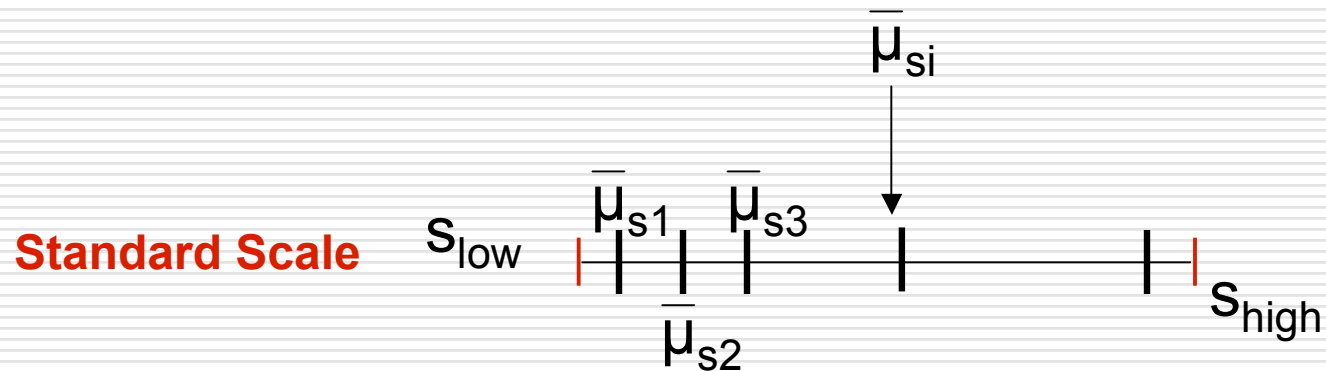


The Intensity Normalization (Training): mapping to Standard scale

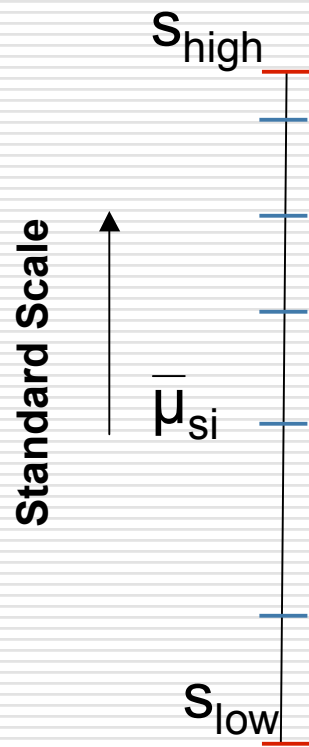


The Intensity Normalization (Training):

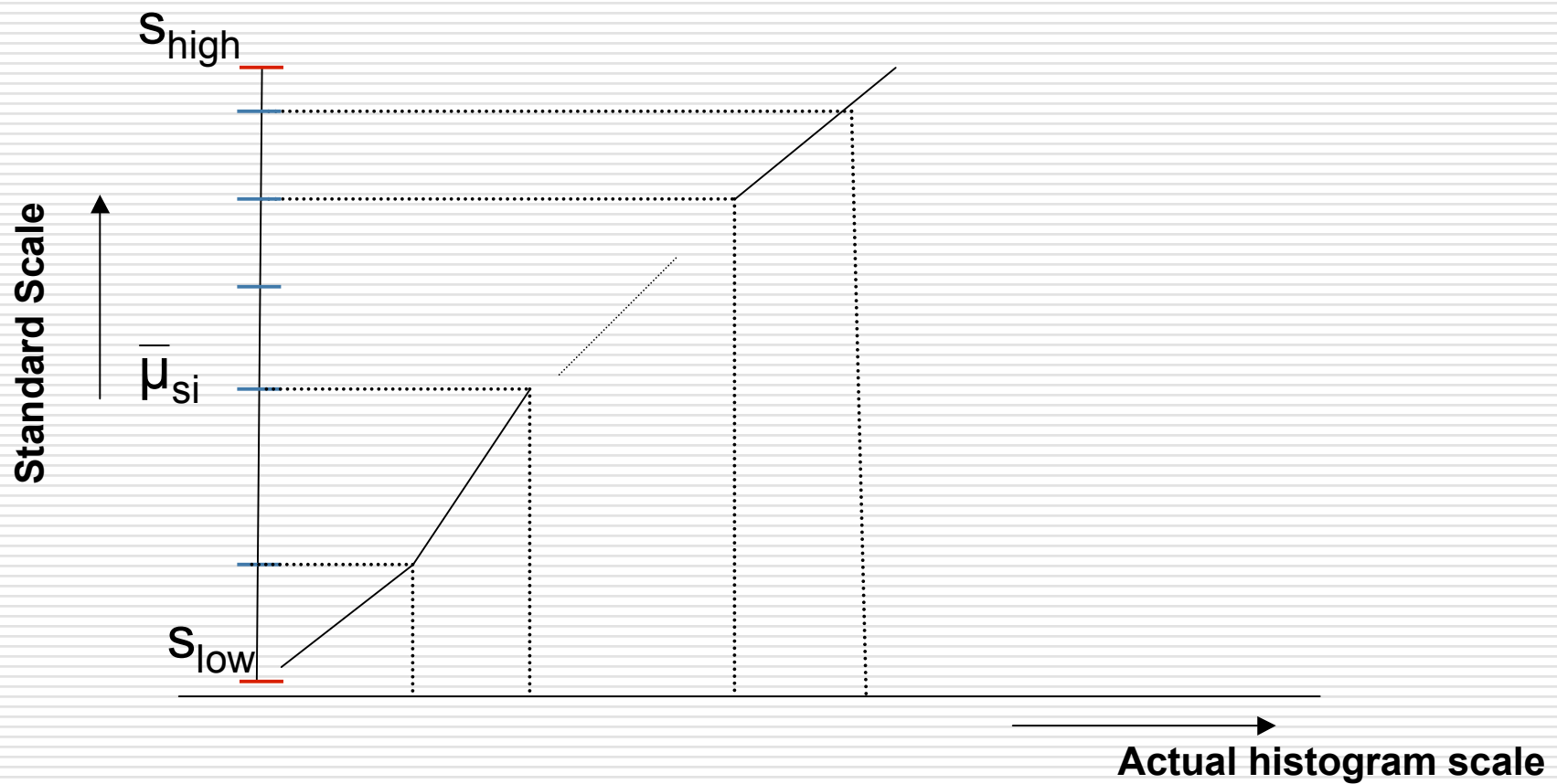
Standard scale



The Intensity Normalization: Standard Scale



The Intensity Normalization: Transformation





Evaluation of Nyul approach

Data Acquisition (data courtesy NeuroRx Research, Montreal)

- 21 multi-spectral (T1w, T2w and PDw) MRI scans
 - 7 subjects each from GE, Philips and Siemens
- Within each sub-group, patients with
 - Varying ventricle sizes
 - Varying MS lesion loads
- Protocols standardized to obtain similar contrasts
- Each MRI volume with
 - 1x1x3 mm resolution
 - 50 slices (vertex to the foramen magnum)

Pre-processing

- Image Modalities' Alignment
 - Image modalities aligned to a common stereotaxic space for deterministic spatial voxel mapping

- Intensity in-homogeneity correction
 - To account for scanner-specific variations

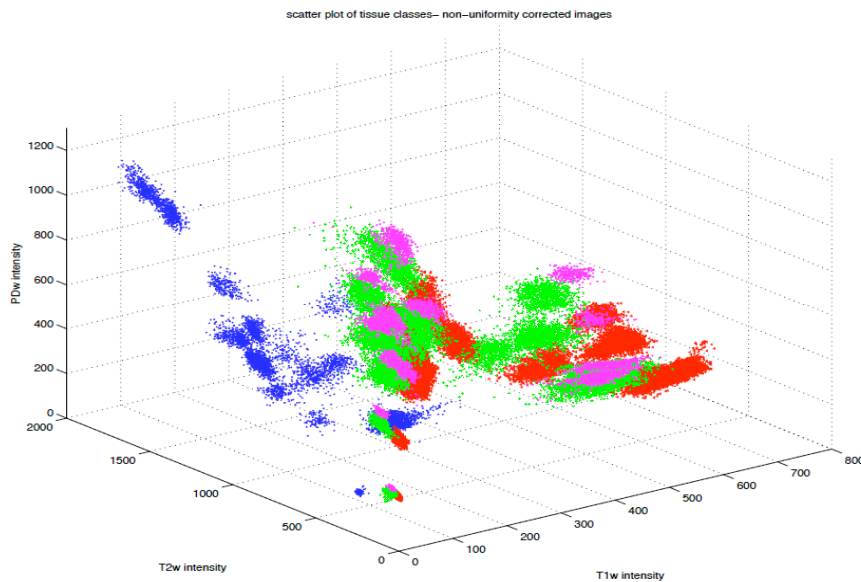
Tissue Sampling

- “Pure” tissue samples
- Tissue types
 - Cortical Gray Matter (CGM), Deep Gray Matter (DGM), White Matter (WM), and Cerebrospinal Fluid (CSF)
- 1000 manual samples for each tissue per MRI
 - about 20 slices (alternate) per brain volume
- Samples obtained both before and after normalization
- Lesions identified automatically followed by expert validations by 5 expert radiologists (consensus)

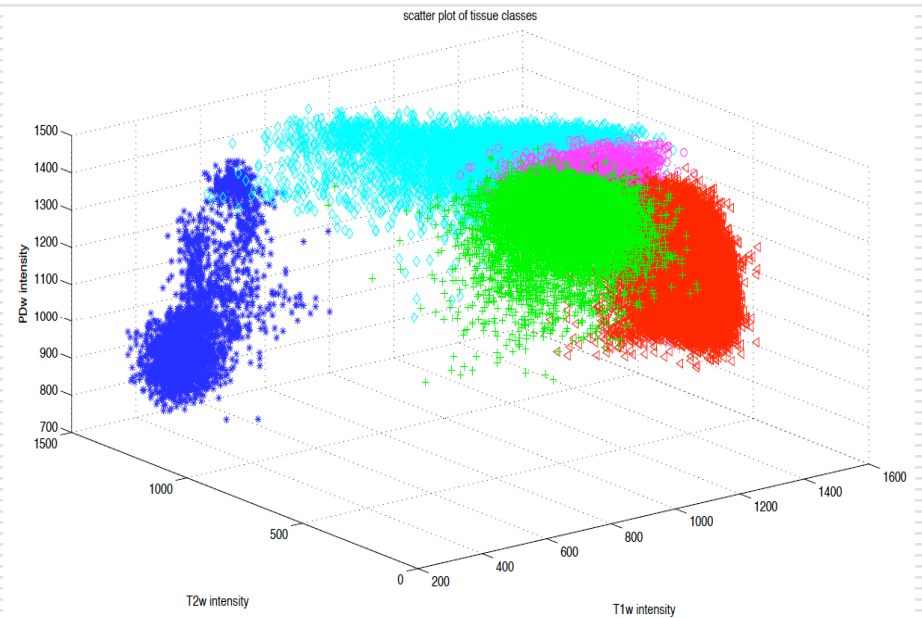
Evaluation

- Training
 - 100 volumes with varying lesion loads and MS pathology
 - Standard scale histogram parameters obtained
 - Same for all modalities
- Standard scale then used in the Transformation stage
- Evaluation Criteria
 - Effect of normalization on heterogenous MRIs
 - Multi-site, Multi-scanner (different manufacturers/brands)
 - Distributional assumption
 - Role of normalization in tissue separation
 - Effect of pathology

Qualitative Evaluation



Un-normalized



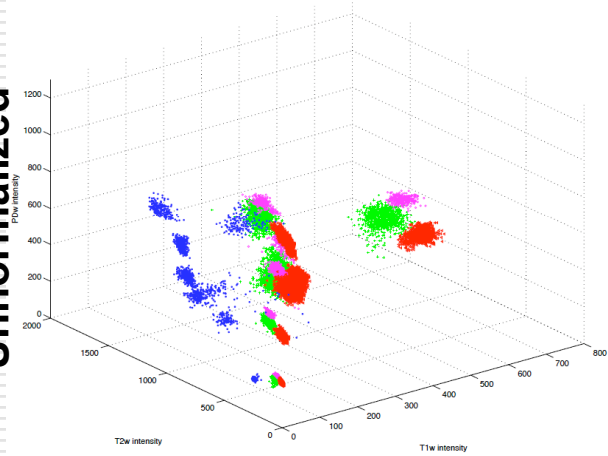
Normalized

Red: WM, Green: CGM, Magenta: DGM, Blue: CSF, Cyan:Lesions

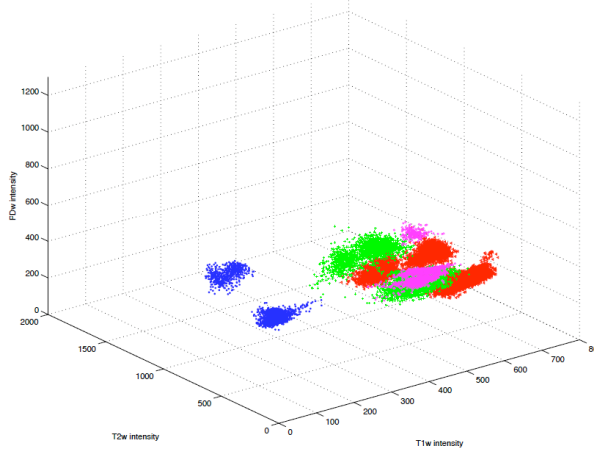
Qualitative Evaluation: by manufacturers

Unnormalized

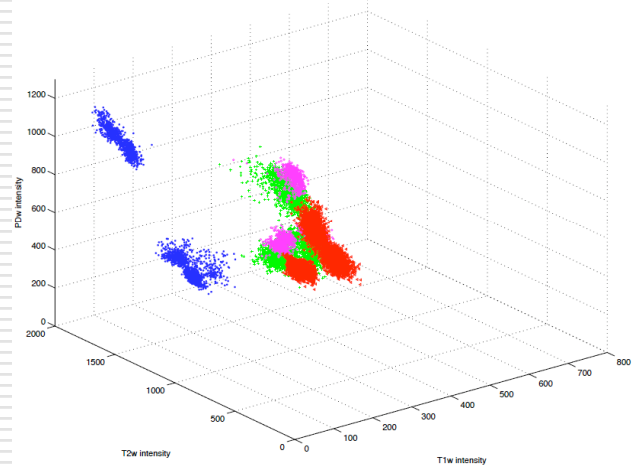
scatter plot of tissue classes GE machines – non-uniformity corrected images



scatter plot of tissue classes, Philips machines – non-uniformity corrected images

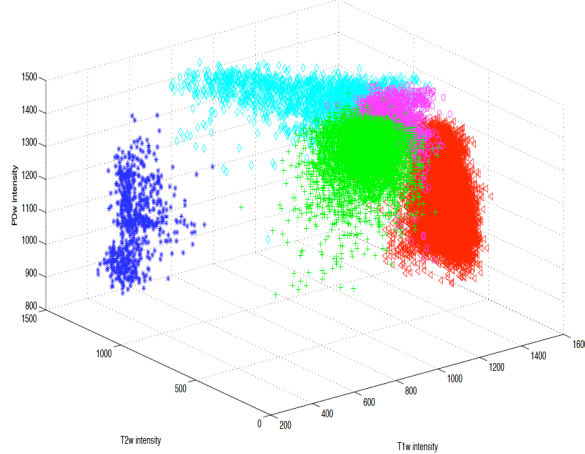


scatter plot of tissue classes, Siemens machines – non-uniformity corrected images



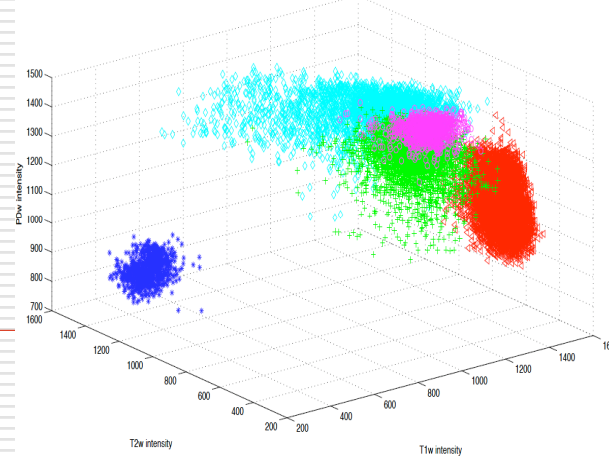
GE

scatter plot of tissue classes GE machines



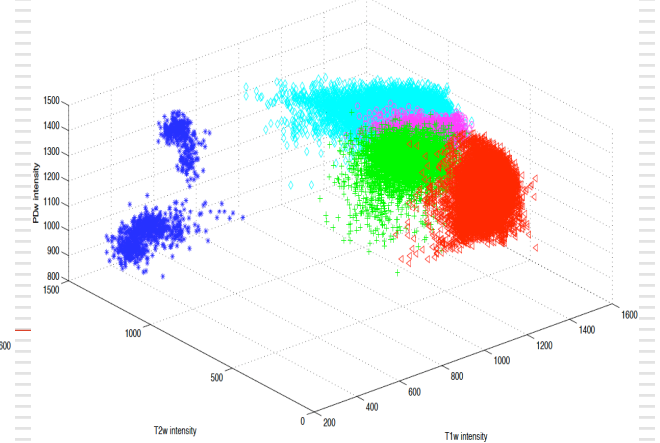
Philips

scatter plot of tissue classes Philips machines



Siemens

scatter plot of tissue classes Siemens machines



Normalized

Quantitative Evaluation

- Tissue behavior in individual modalities
- Effect on Gaussian distribution assumption
- Jeffrey Divergence criterion

$$JD(p, q) = \int_{-\infty}^{\infty} p(x) \log \frac{p(x)}{q(x)} dx + \int_{-\infty}^{\infty} q(x) \log \frac{q(x)}{p(x)} dx$$

Verifying Gaussian assumptions

- For un-normalized images:
 - Calculate intensity histogram for each tissue type from the input image (bin size 100). Call it **D1-u**
 - Calculate data mean m and covariance c
 - Generate a Gaussian at (m, c) . Call it **D2-u**
 - Calculate the Jeffrey Divergence between D1-u and D2-u: **JD-u**

- For normalized images
 - Do as above and calculate **JD-n**

- Compare JD-u and JD-n

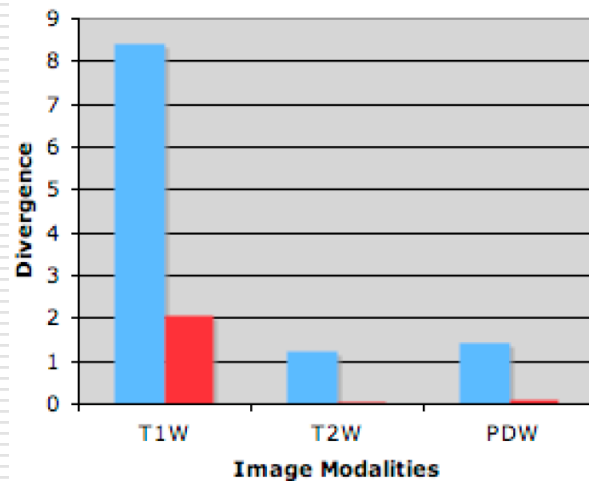
Hypothesis for Evaluation

- ☐ If the data comes from a Gaussian distribution then JD-u shouldn't be large
- ☐ Further if normalization doesn't have any effect on data distribution then the difference in JD-u and JD-n should be insignificant

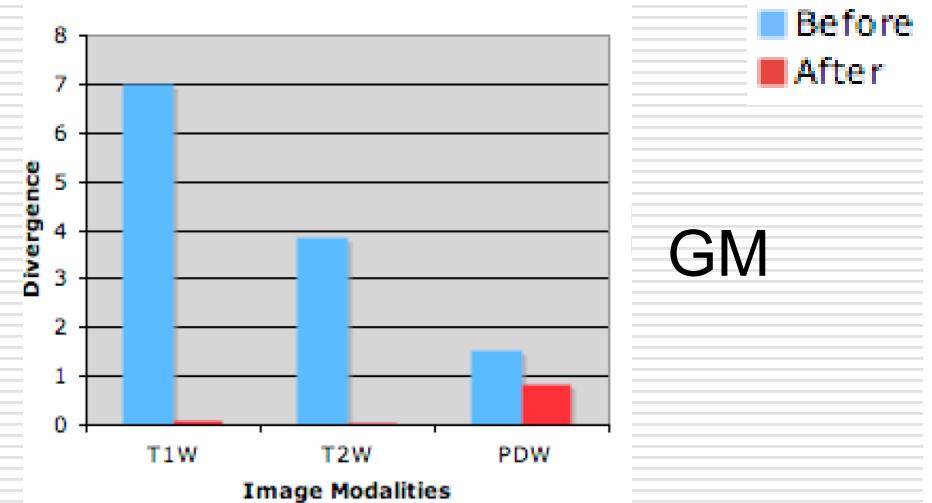
Results

Jeffrey Divergences: All machines

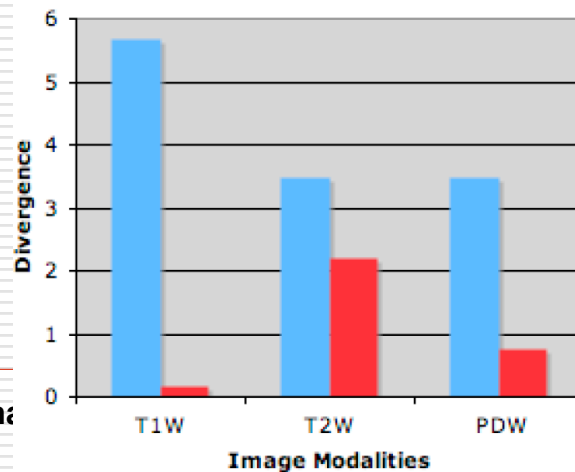
WM



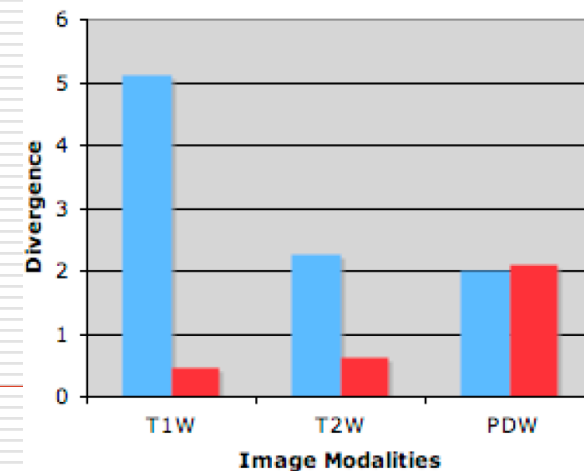
GM



CSF



Lesions



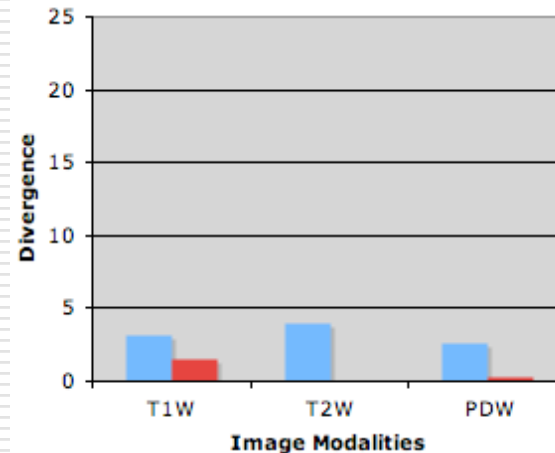
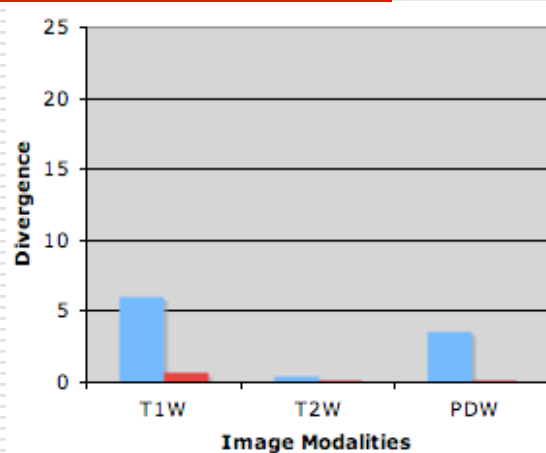
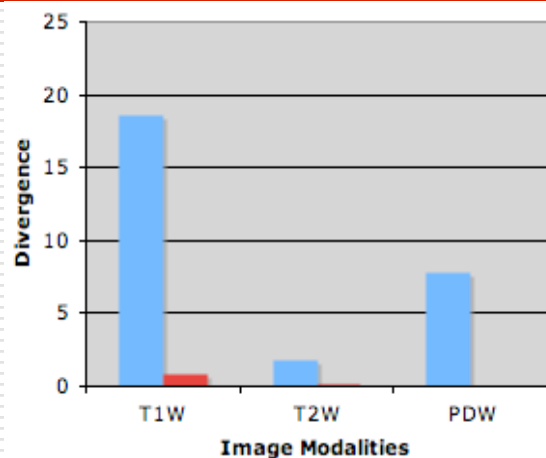
Moham

IAMS-2009

Results

WM and GM : Jeffrey Divergences, by Manufacturers

WM

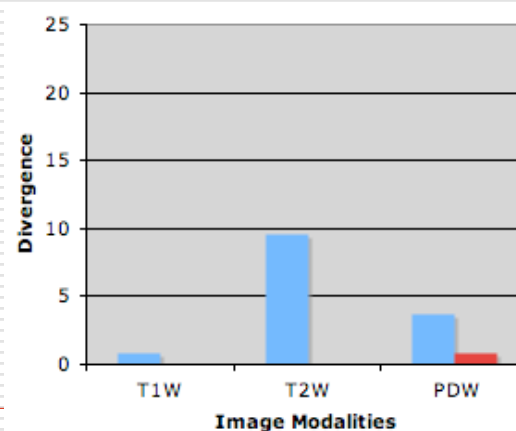
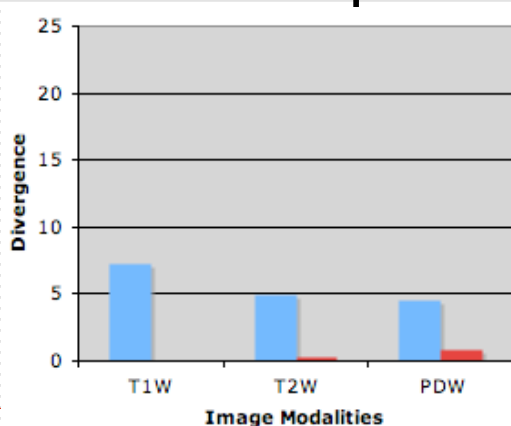
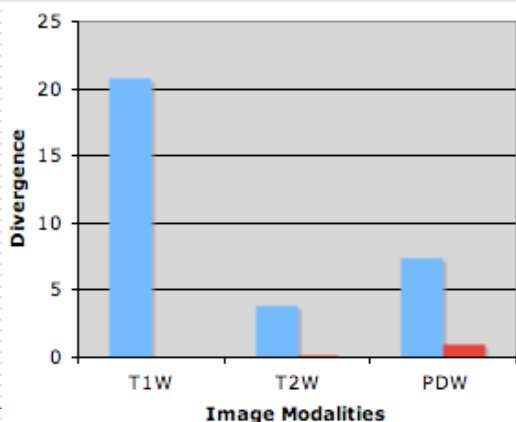


GE

Philips

Siemens

GM

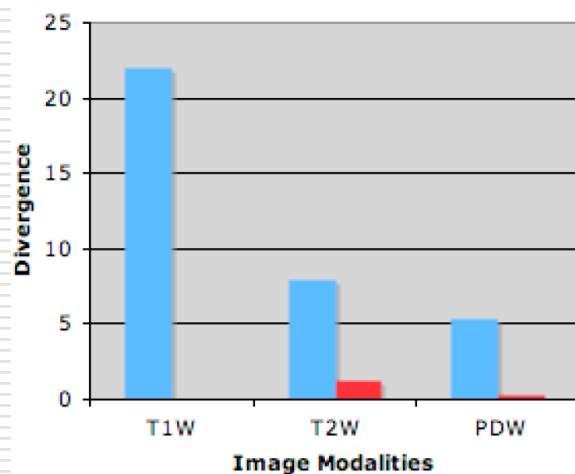


Mohak Shah

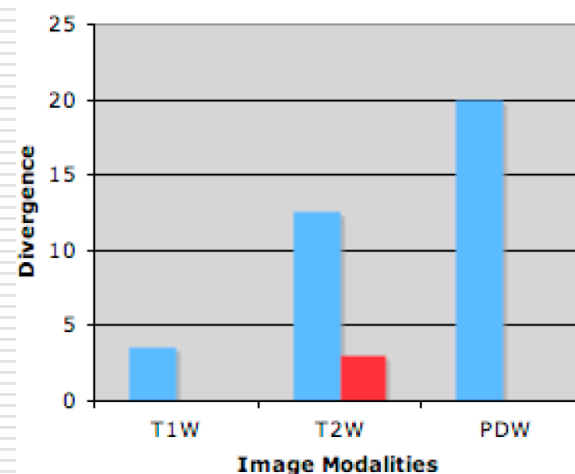
MIAMS-2009

Results

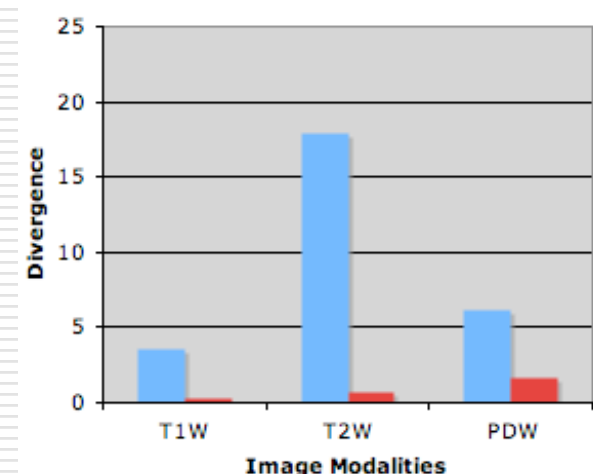
CSF : Jeffrey Divergences, by Manufacturers



GE



Philips

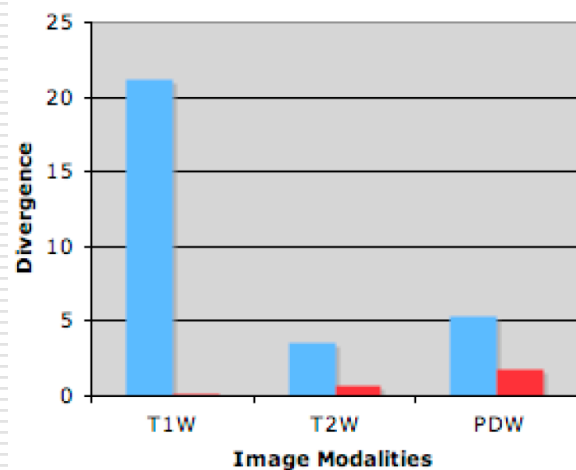


Siemens

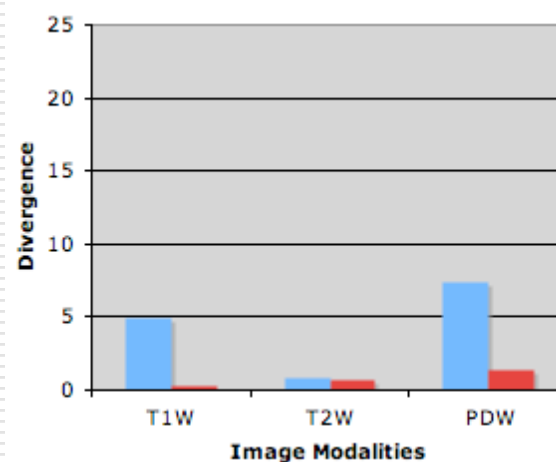
■ Before
■ After

Results

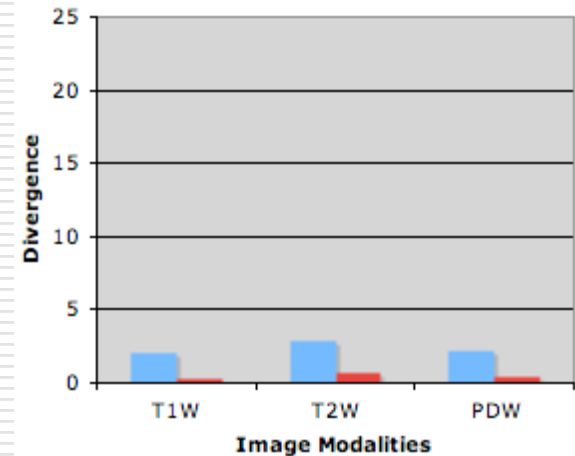
Lesions : Jeffrey Divergences, by Manufacturers



GE



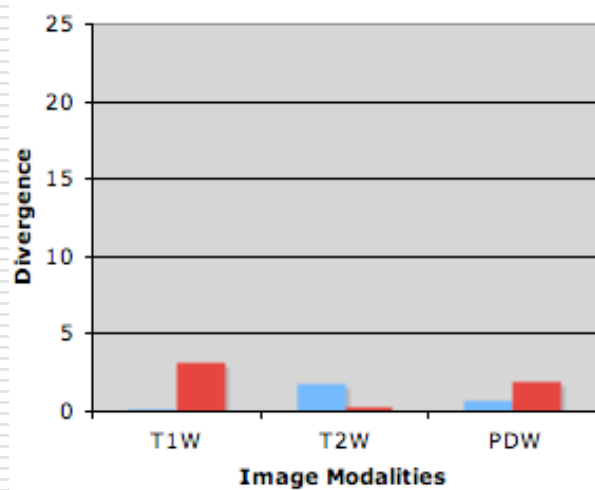
Philips



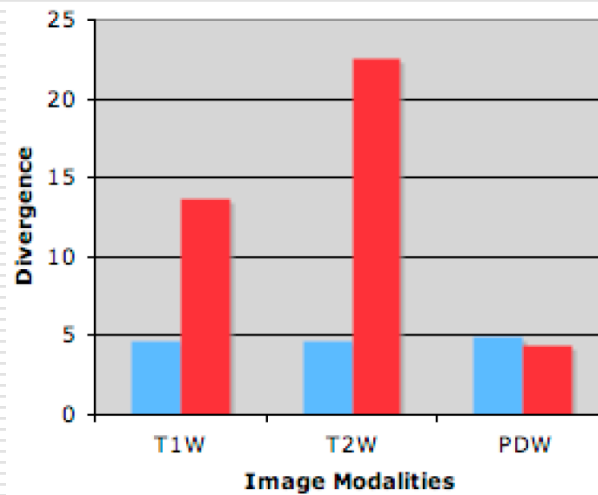
Siemens

■ Before
■ After

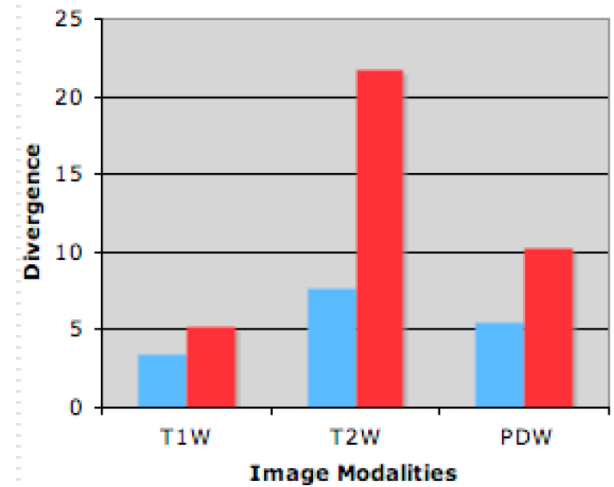
Tissue Contrast



WM:GM



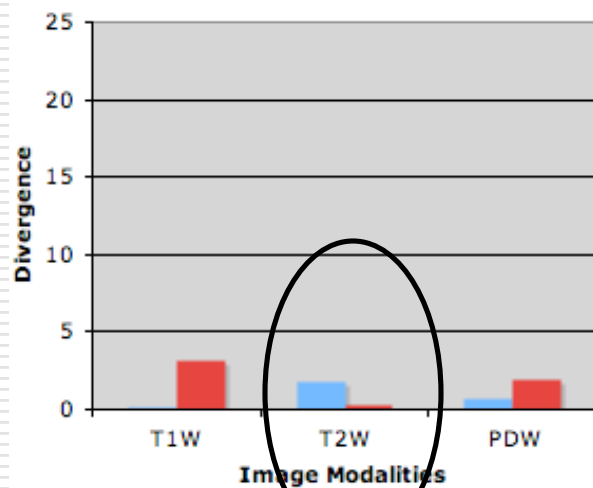
WM:CSF



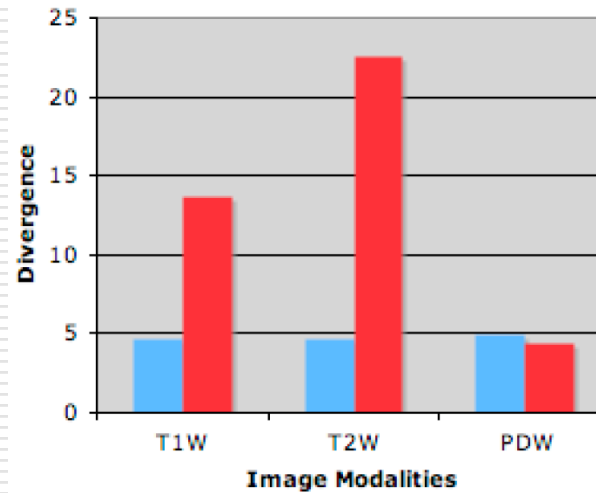
GM:CSF

Before
After

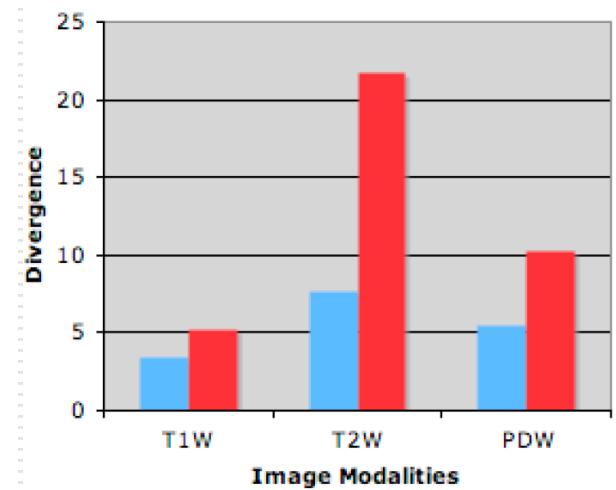
Tissue Contrast



WM:GM



WM:CSF



GM:CSF

Conclusion and Future Work

- Nyul approach affects the tissue intensity distribution
 - Homogenizing same tissues
 - Better separation between different tissues
- Makes the distribution amenable to application of segmentation algorithms
- Robust in the presence of pathology
- **To Do:**
 - Effect on segmentation results
 - Lesion load dependent analysis
 - Incorporating this knowledge into automatic approaches



Thank You