

MRI and MRSI Application in Neurodegeneration

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Magnetic Resonance Imaging(MRI) MR Spectroscopic Imaging(MRSI)

I. H-1 Magnetic Resonance Spectroscopic Imaging in Alzheimer's Disease, Multiple Sclerosis and Epilepsy

II. Micro Na-23 MRI and micro F-18 FDG-PET of cancer and apoptosis

Neurodegeneration: AD, Epilepsy, MS

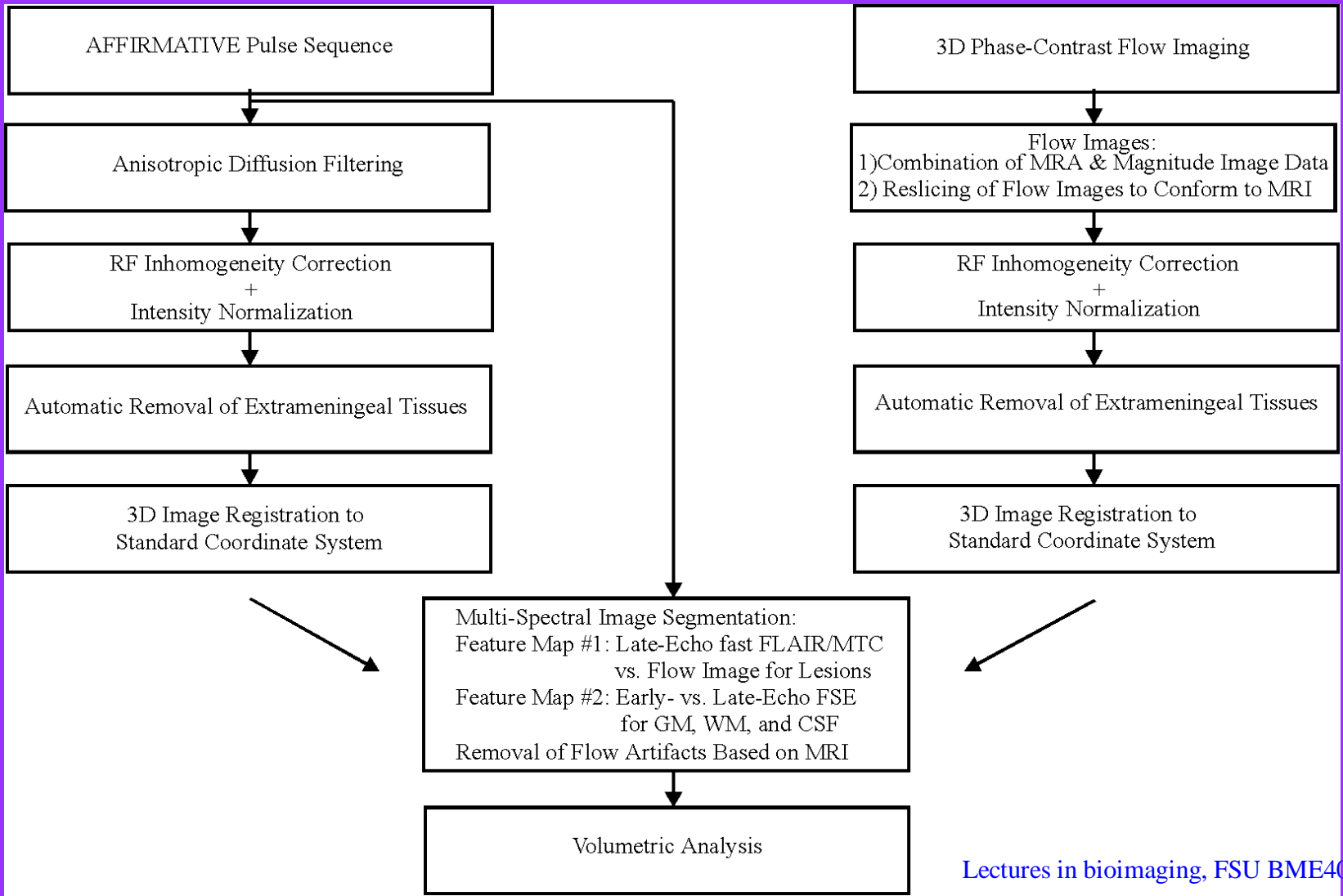
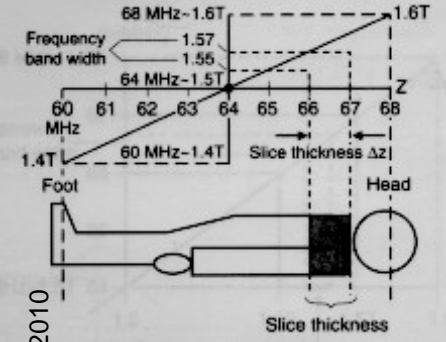
Hypothesis

**MRI combined with MRSI characterize &
define the disease better**

MRI Imaging: Concepts

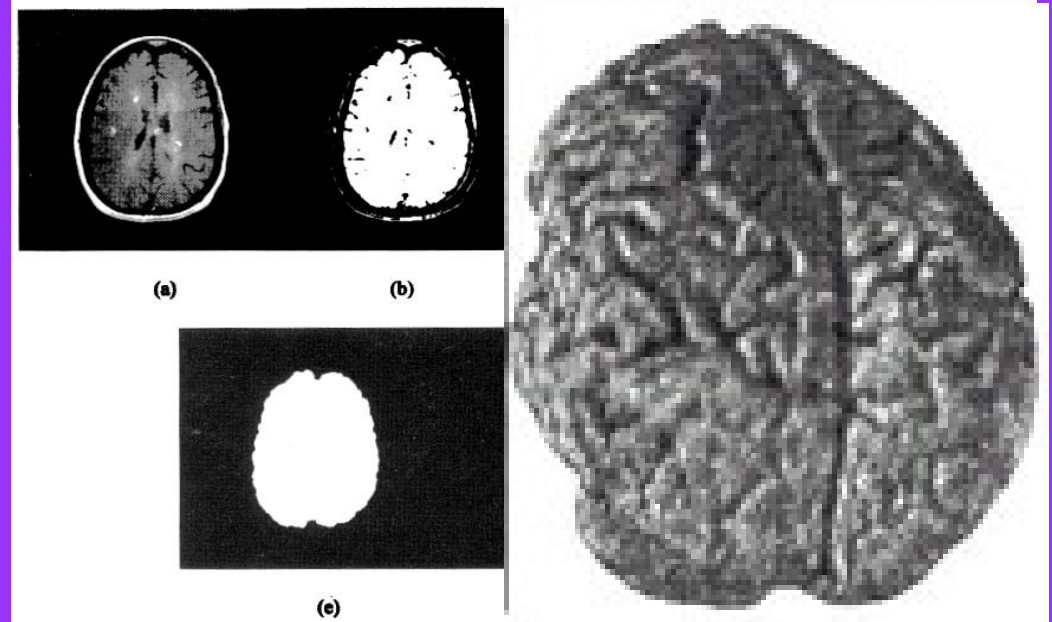
- **Proton spins resonate with RF pulse in high magnetic field: Slice imaging**
- **RF pulse sequences: Spin-echo, Inversion recovery, Gradient echo, phase contrast, Time-of-flight(TOF) sequences**
- **T1, T2, proton density weighting by TE, TR, TI etc.**
- **Data acquisition: 2D slice, 3D volume imaging.**
$$S(k_x, k_y, t) = \iint \Sigma \rho \cdot e^{\gamma \Delta B_0 \gamma \phi(r) + (TI-TR/T1)}$$
- **Surface Rendering and Volume Rendering**
- **Manipulation and Analysis: Interactive MRI and MR Spectroscopic Imaging**

1.5 T MRI is Enough to Generate Brain Image

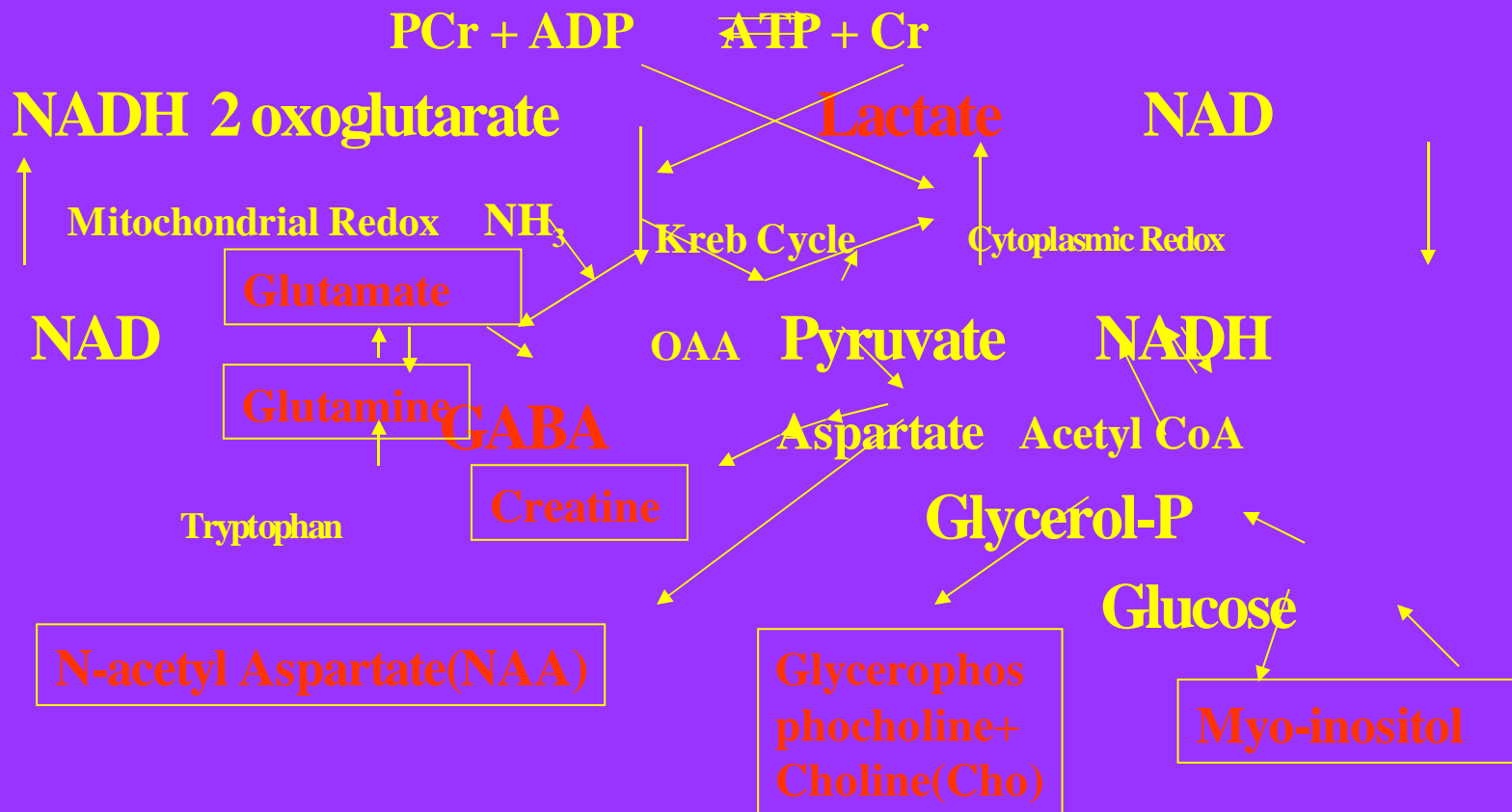


Brain Tissue Composition

- **Cortex**
- **Sub-cortical**
- **Cingulate**
- **Ventricles**
- **Gyri and Sulci**
- **Frontal, Parietal, Temporal, Inter-hemispheric Fissure**
- **Distribution of gray matter, white matter and CSF as total content $\rho=(GM+WM+CSF)$**



MR visible metabolites in brain



Metabolite signal = (MRI resolution x SRF x chemical shift profile) x H-1 density (MRSI resolution)

Proton Magnetic Resonance Spectroscopic Imaging(MRSI) Technique

- **Data Acquisition:**

- H-1 Multi Slice SI(MSSE) sequence-**Metabolite maps**

- PE 36 diameter, FOV 280.280, (TE 25 TR 1800 TI 170)ms

- Stimulated (CHESS) SI sequence (STEAM)-Spectral map**

- PE 32 x 32, TR/TE 1000/30, NEX 2, VOI 240 mm², ROI .75 x .75 x 1.5 cm³

- Point-Resolved SI sequence (PRESS)-Single Spectrum**

- PE 24.24, FOV 210.210, (TE 25 TR 1800)ms, voxel 2.2 ml

- Outer volume suppression for minimizing extrameningial lipids

- Variable TR for reducing scan time

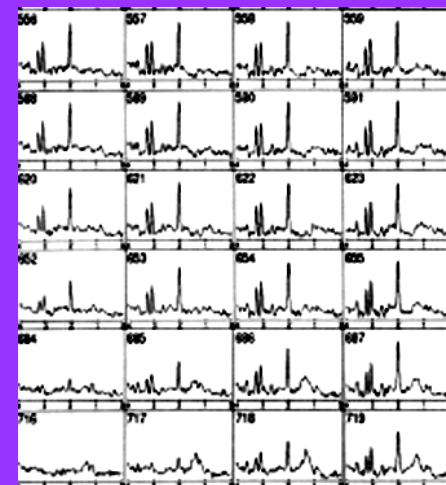
- **Image Postprocessing**

- Segmentation, Co-registration (**MRIAP, Viewer**)

- Automatic Spectral Image Analysis software(**APSIP**)

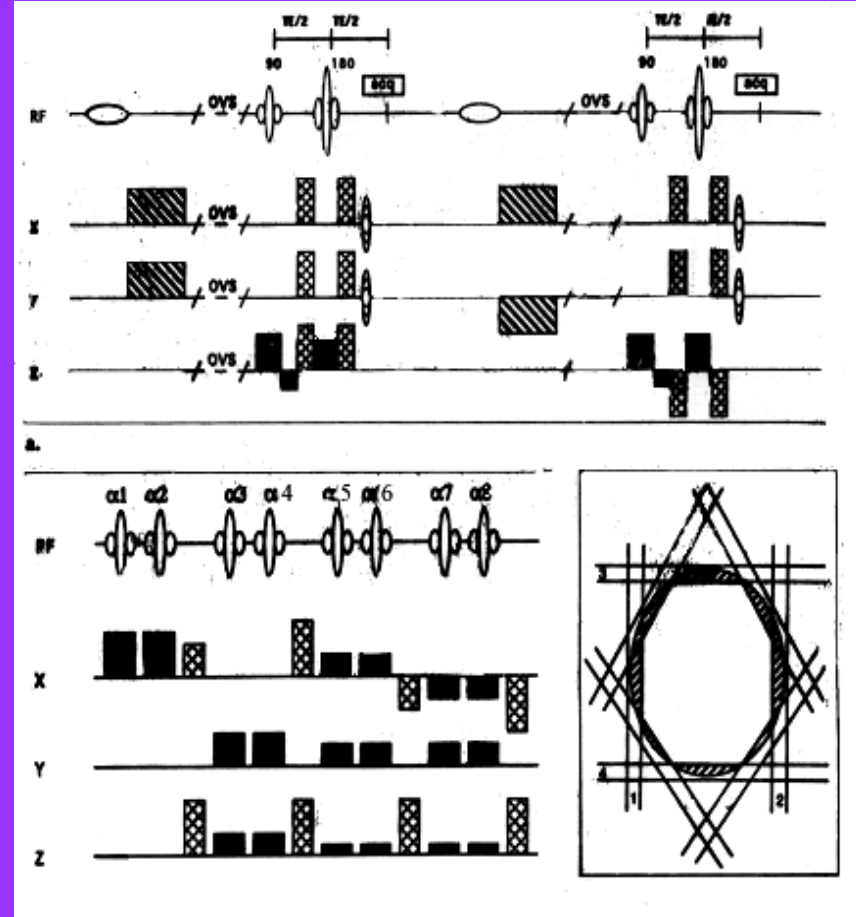
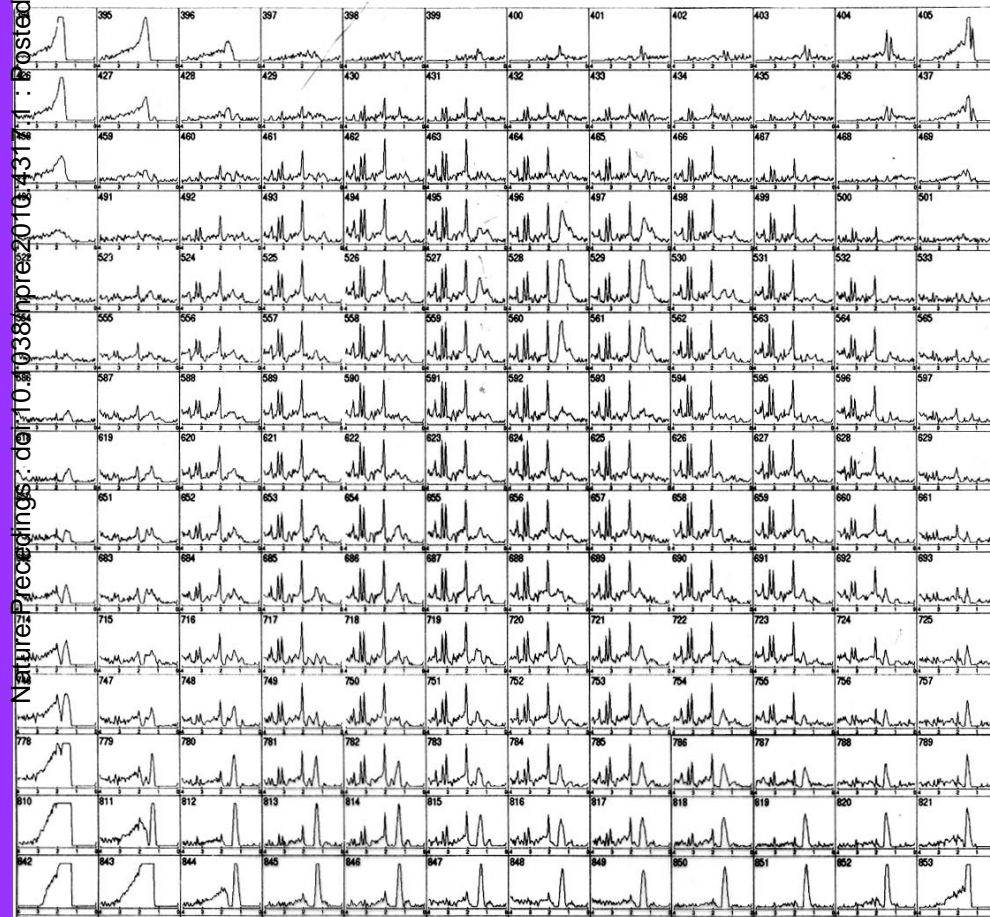
- Metabolic map generation, Spectral analysis

- Interfacing MRSI with MRI (Co-analysis by **SID**)



Multi-slice octagonal VOI: STEAM Chemical Shift Imaging

Nature Precedings doi:10.1038/npre201043171

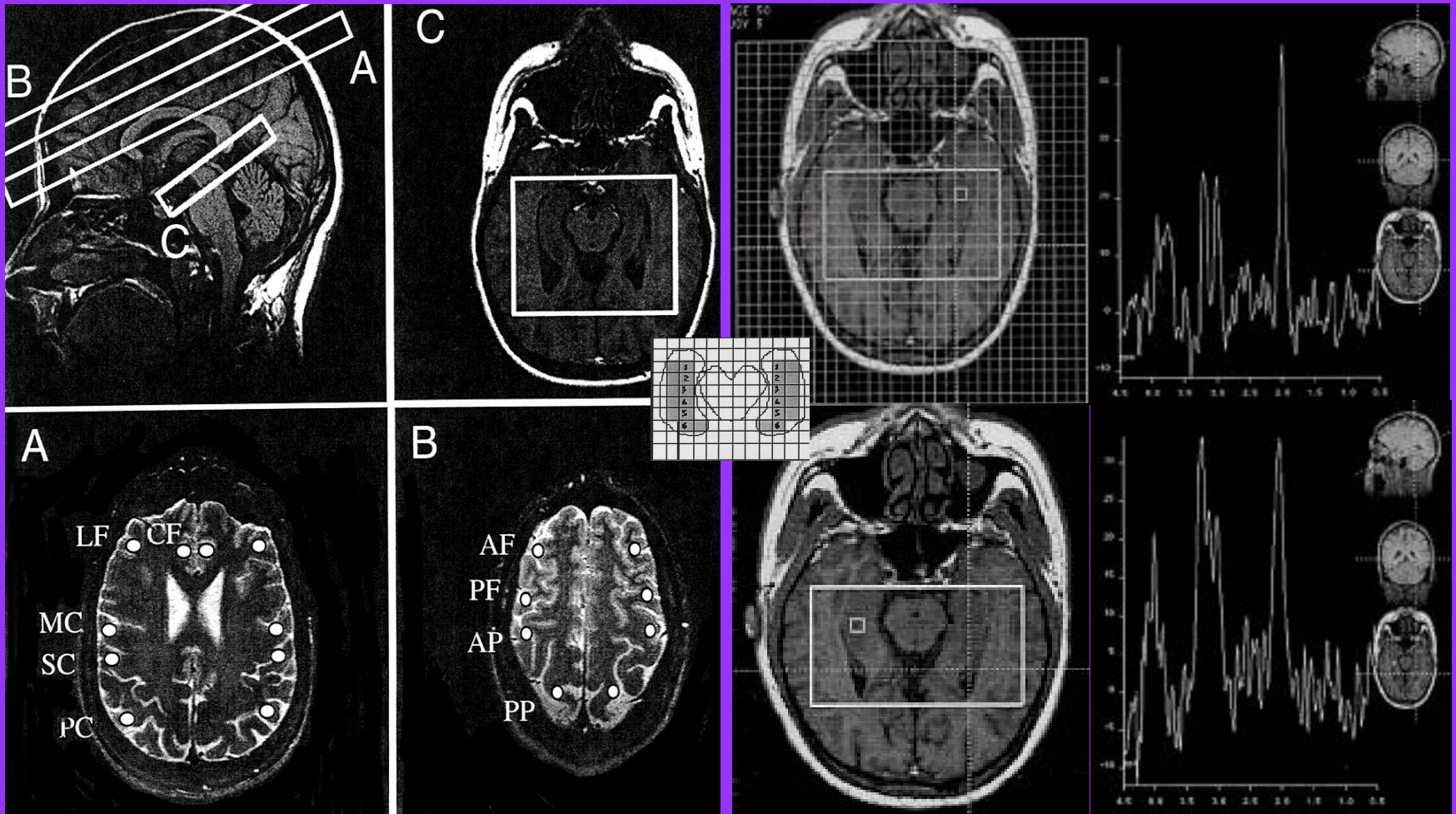


MRI+MRSI in Alzheimer's Disease (AD) and STVD


1. **AD is diffused cortical neurodegenerative disease**
 - **Memory Loss is associated with Reduced NAA(Neuronal Loss) and Enhanced Choline (Inflammation,demyelination)**
 - **Subcortical Ischemic Vascular Dementia**
 - **MRI + MRSI define it better**
 - **MRI offers visible hippocamcal changes (segmentation)**
 - **MRSI offers localized neurochemical changes for lateralization, test for asymmetry (co-analysis)**

H-1 MRSI + MRI predict Hippocampal Volumes and Neurometabolites in Alzheimer's Disease

Nature Precedings : doi:10.1038/npre.2010.4317.1 : Posted 28 Mar 2010



MRI and MRSI Image Post Processing and Analysis in AD vs SIVD

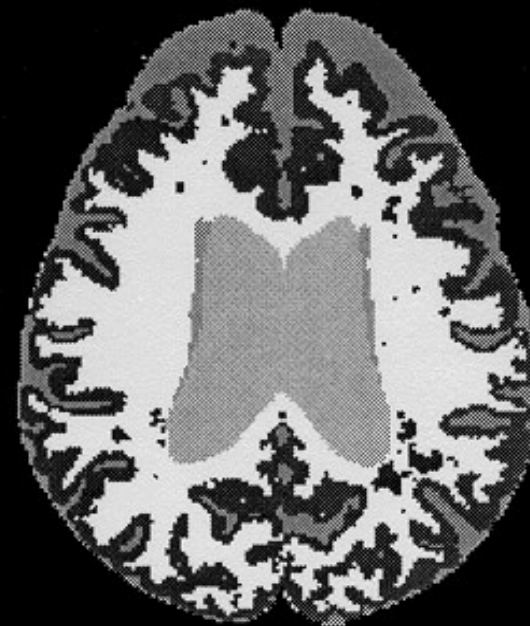
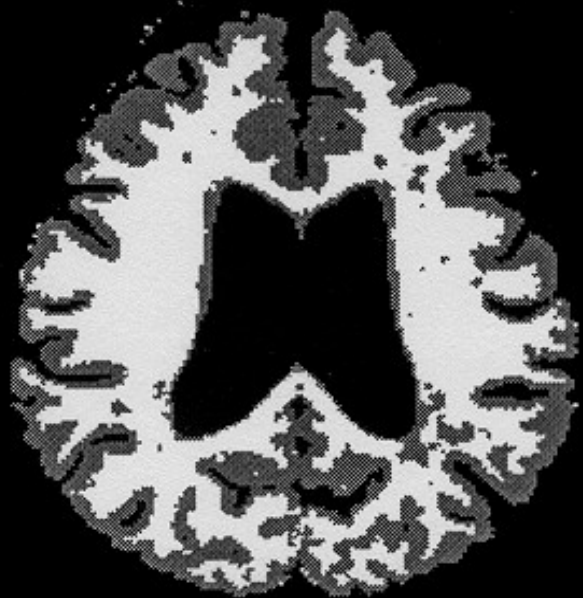
- **Image Post-Processing:** MRI-Ventricular dilatation, Sulcus widening, subcortical WMSH
 - MSSI and CSI-Zero-filling, exponential line broadening(time domain), Gaussian Multiplication(spatial domain), Water suppression(digital filter+linear interpolation in FID), fat removal,FFT,Atrophy Correction
 - Automated Spectral Analysis: Simulated parametric peak amplitude,phase,frequency(sinusoidal LG shape), FFT+Baseline Wavelet-LM Optimization Iterations Curve fits and metabolite concentrations
 - **Statistics:** 
1. WM and GM(Frontal,Parietal) NAA/Cr,NAA/Cho, Cho/Cr): Control vs AD vs SIVD
 2. Ventricular Dilatation,WMSH,Sulcus Widening, Hippocampal Volume: Control vs AD vs SIVD
 3. Hippocampal Volume, NAA/Cr, NAA/Cho, Cho/Cr: Control vs AD vs SIVD
 4. Total(GM+WM) Metabolite Differences: AD vs Control and AD vs SIVD

MRI segmentation(1st step)

- **Removal of extrameningial tissue and scalp**
- **Co-registration of DSE images with interleaved images(Wood Algorithm)**
- **3D inhomogeneity correction**
- **Segmentation by K-means cluster analysis - tissue-seeds defined around pixel intensity histograms**
- **Automated edge detection for hippocampus area and hippocampus voluming**

Tissue MRI segmentation (supervised K-NN cluster method)

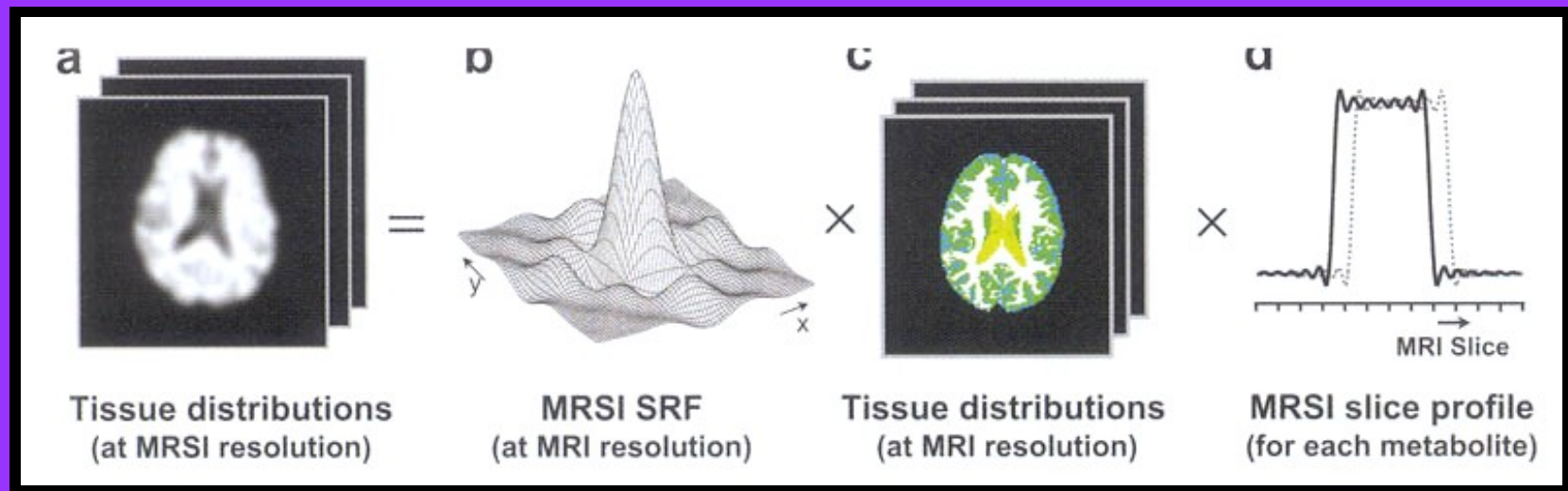
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MRSI post-processing(2nd step)

- Parametric Automated spectral peak editing
- *a priori* information of metabolites and voxel-by-voxel training data for tissue composition(CSF nulling)
- Point Spread Function and Chemical shift displacement(**MRSI Resolution**)
- Data Processing in the X, Y co-ordinates
- Co-registration and segmentation for serial scans

Resolution at MRI and MRSI slices: Multi-slice MR Spectroscopic Imaging Display(SID)

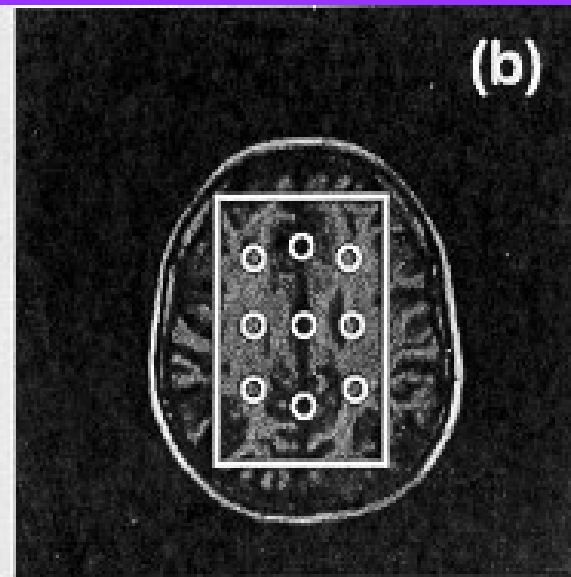
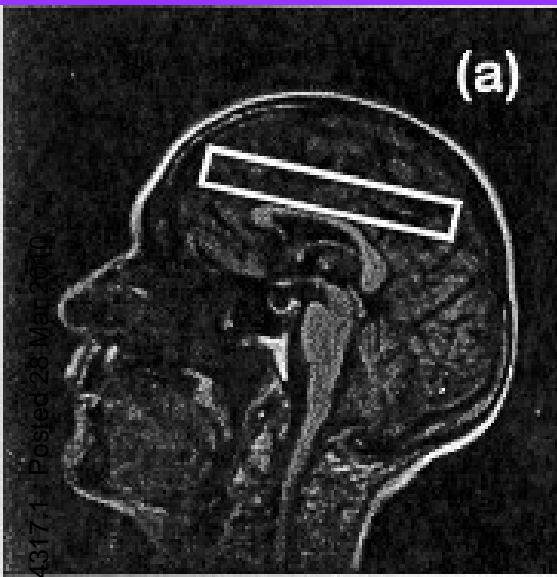


Effective tissue contribution in spectroscopic volume =

SRF x tissue distribution(MRI resolution) x MRSI Slice selective profile

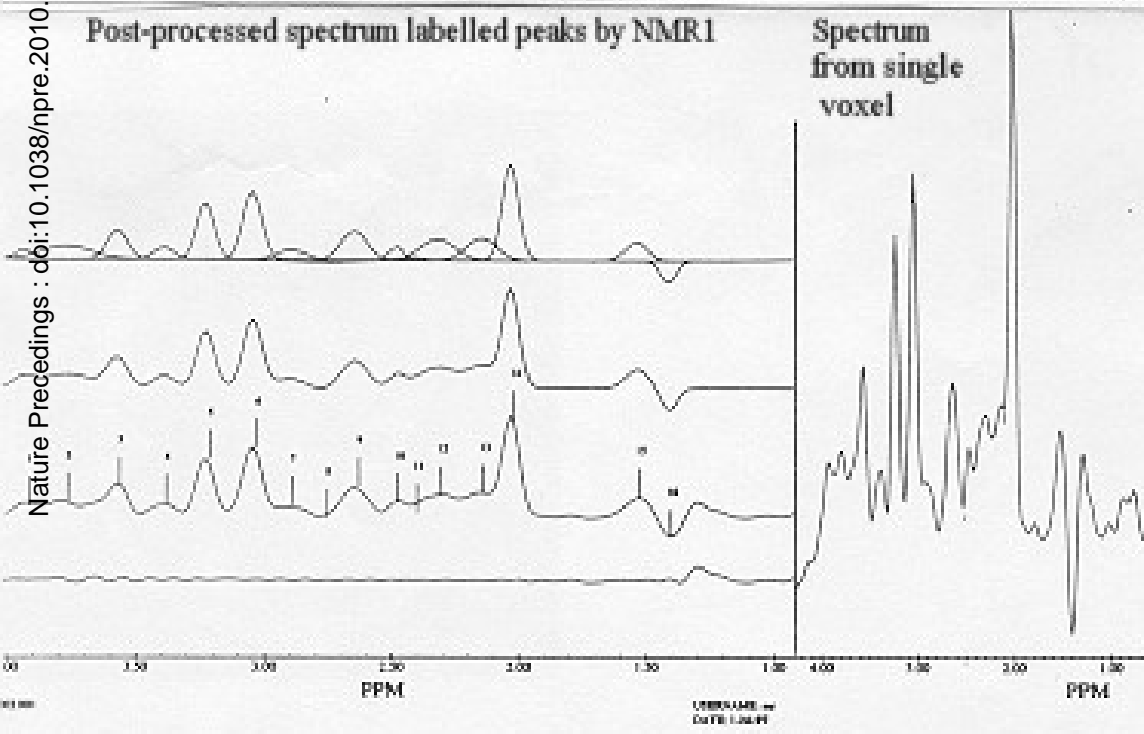
MRSI: Spectra Post-processing (NMR-1 program) (Step 3)

- **Deconvolution**
- **Gaussian apodization**
- **Zero-filling to matrix 32 x 32 x 1024 points**
- **Fourier Transformation**
- **Phase and baseline correction**
- **Peak editing and reference with phantom**
- **Iteration and curve-fitting**



1.5 T MR Spectroscopy and Image Processing of Alzheimer's Disease

- (a) Sagittal scout image
- (b) Location of ROI voxels



- Lower-left: Curve fitted spectrum
- Lower-right: Metabolite peaks

MRI segmentation and co-analysis with H-1 MRSI (SID)(step 4)

- Tissue content $\rho = \text{GM} + \text{WM}$ (from MRI)
- $f = \text{GM} / (\text{GM} + \text{WM})$
- Co-registration of MRI with MRSI data
- Metabolite intensity correction for NAA, Cr and Cho as: $\text{NAA}^{\text{corrected}} = \text{NAA} / \rho$ (MRSI)
- CSF signal nulling
- 3D inhomogeneity correction by digital filter

Regional Differences of Metabolites in brain: AD

| NAA [mM] | Brain Region | | |
|---------------------|--------------|-------------|-------------|
| | Frontal | Medial | Posterior |
| Gray Matter | | | |
| AD | 8.59 ± 0.4 | 9.35 ± 0.3 | 9.43 ± 0.3 |
| Control | 10.04 ± 0.4 | 9.75 ± 0.4 | 10.52 ± 0.4 |
| % Difference | - 14.4 * | - 4.1 | - 10.3 * |
| White Matter | | | |
| AD | 8.33 ± 0.3 | 9.70 ± 0.3 | 8.94 ± 0.2 |
| Control | 9.25 ± 0.2 | 9.63 ± 0.2 | 9.48 ± 0.3 |
| % Difference | - 10.1 ** | + 0.7 | - 5.7 |
| Cr [mM] | | | |
| Gray Matter | | | |
| AD | 8.52 ± 0.3 | 8.08 ± 0.2 | 8.11 ± 0.3 |
| Control | 8.70 ± 0.3 | 7.29 ± 0.3 | 8.15 ± 0.3 |
| % Difference | - 2.0 | + 10.8 | - 0.5 |
| White Matter | | | |
| AD | 6.39 ± 0.2 | 6.19 ± 0.2 | 6.72 ± 0.2 |
| Control | 6.52 ± 0.3 | 6.12 ± 0.2 | 6.74 ± 0.2 |
| % Difference | - 2.1 | + 1.1 | - 0.3 |
| Cho [mM] | | | |
| Gray Matter | | | |
| AD | 1.85 ± 0.09 | 1.55 ± 0.06 | 1.34 ± 0.07 |
| Control | 1.82 ± 0.07 | 1.56 ± 0.08 | 1.38 ± 0.07 |
| % Difference | + 1.7 | - 0.7 | - 3.2 |
| White Matter | | | |
| AD | 1.61 ± 0.07 | 1.49 ± 0.05 | 1.30 ± 0.05 |
| Control | 1.71 ± 0.08 | 1.55 ± 0.06 | 1.26 ± 0.04 |
| % Difference | - 5.8 | - 4.1 | + 3.1 |

*p < 0.03; ** p < 0.003; both by ANCOVA;

MRI+MRSI Predicts Better Tissue Composition in AD

| | AD | Control | % diff. |
|----------------------------------|------------|------------|---------|
| HP-volume(mm³) | | | |
| Right | 1982 ± 134 | 2884 ± 102 | 31.1 |
| Left | 1868 ± 88 | 2943 ± 86 | 36.5 |
| Ventricular CSF(%) | 4.2 ± 0.3 | 2.8 ± 0.3 | 33.3 |
| Sulcus CSF (%) | 23.4 ± 2 | 18.2 ± 0.5 | 22.2 |
| White Matter(%) | 35.2 ± 0.9 | 38.1 ± 0.8 | 7.6 |
| Cortical GM(%) | 38.8 ± 1.1 | 42.2 ± 0.6 | 8.0 |
| Subcortical GM(%) | 1.2 ± 0.08 | 1.4 ± 0.03 | n.s |
| TIV(cm³) | 1342 ± 5 | 1402 ± 52 | n.s. |

MRSI: Metabolite* distribution and tissue content in AD

| Metabolite | AD | Control | %Diff | | AD | Control | % Diff |
|----------------|------------|-------------|-------|-------------------------------|-------------|-------------|--------|
| NAA(mM) | | | | NAA/Cr | | | |
| Right | 7.55 ± 0.5 | 10.01 ± 0.6 | 13.2 | Right | 1.08 ± 0.06 | 1.39 ± 0.03 | 20.8 |
| Left | 7.61 ± 0.4 | 9.82 ± 0.9 | 22.6 | Left | 0.96 ± 0.04 | 1.31 ± 0.02 | 26.7 |
| Cho(mM) | | | | NAA/Cho | | | |
| Right | 2.02 ± 0.7 | 2.08 ± 0.2 | 2.9 | Right | 3.74 ± 0.07 | 4.81 ± 0.09 | 22.2 |
| Left | 2.04 ± 0.4 | 1.89 ± 0.5 | 7.4 | Left | 3.73 ± 0.03 | 5.1 ± 0.06 | 26.9 |
| Cr(mM) | | | | ⊗ Tissue content ρ (%) | | | |
| Right | 7.02 ± 0.6 | 7.75 ± 0.5 | 9.4 | Right | 84 ± 3 | 98 ± 2 | 14.3 |
| Left | 7.96 ± 0.6 | 7.49 ± 0.8 | 5.9 | Left | 87 ± 3 | 96 ± 3 | 0.93 |
| | | | | Gray matter index (f) | | | |
| | | | | Right | 0.45 ± 0.03 | 0.55 ± 0.05 | 18.2 |
| | | | | Left | 0.62 ± 0.02 | 0.62 ± 0.04 | n.s. |

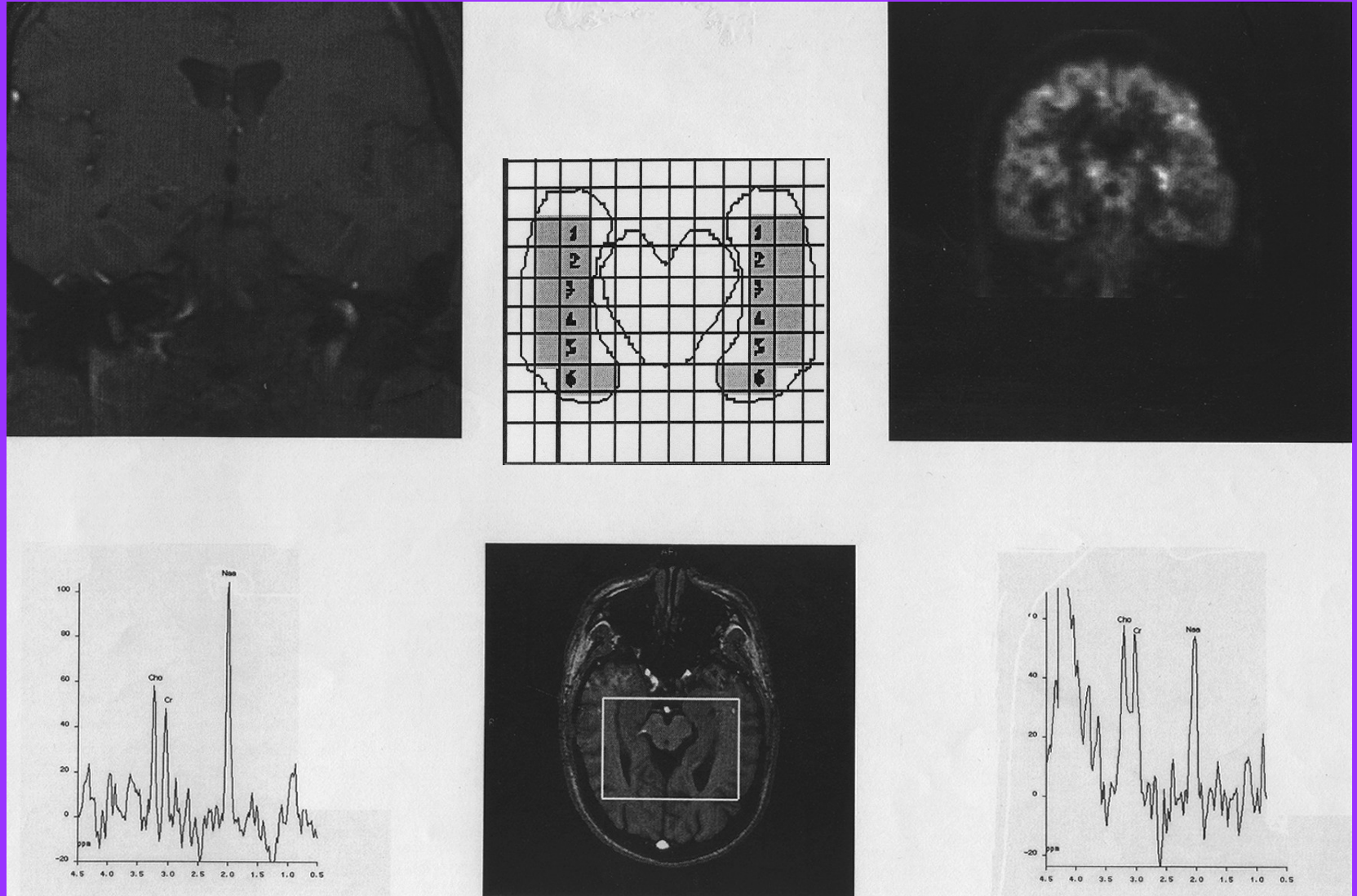
*Atrophy corrected metabolite concentrations of NAA, Cho, and Cr and NAA/Cr and NAA/Cho ratios from right and left hippocampus in AD patients and control subjects.

⊗Tissue content ρ (in percent of the MRSI voxel volume) and gray matter index f of the MRSI voxels positioned at right and left hippocampus, characterizing MRSI partial volume effects.

AD: Lateralization and Asymmetry

| AD | Concordant | Discordant | non-lateralized |
|---------------------------|------------|------------|-----------------|
| FDG-PET | | | |
| Asymmetry Index > 3 | 12 | 0 | 1 |
| Hippocampus Volume | | | |
| AI > 8 % | 10 | 0 | 3 |
| T2 | | | |
| AI > 4 % | 9 | 0 | 4 |
| NAA/(Cho+Cr) | | | |
| AI > 12 % | 7 | 1 | 5 |
| NAA | | | |
| AI > 12 % | 7 | 0 | 6 |

Lateralization of Hippocampus in AD: Reduced NAA and High Choline Peaks

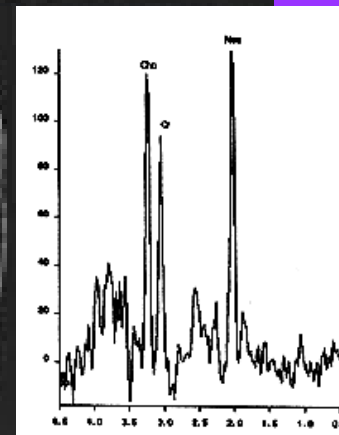
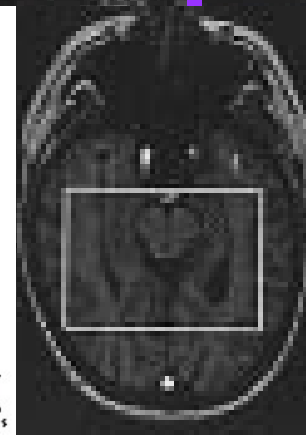
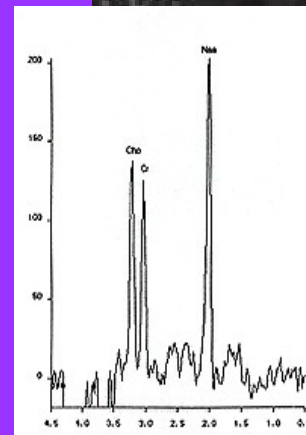
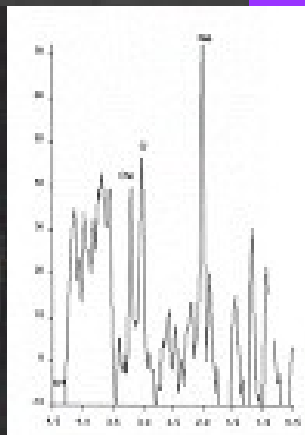
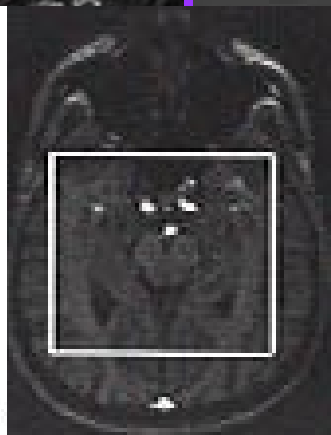
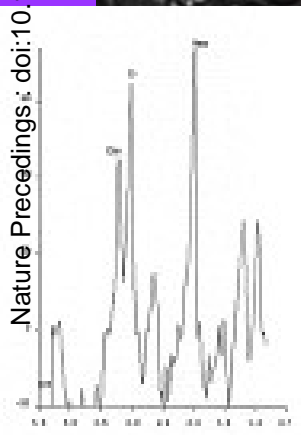
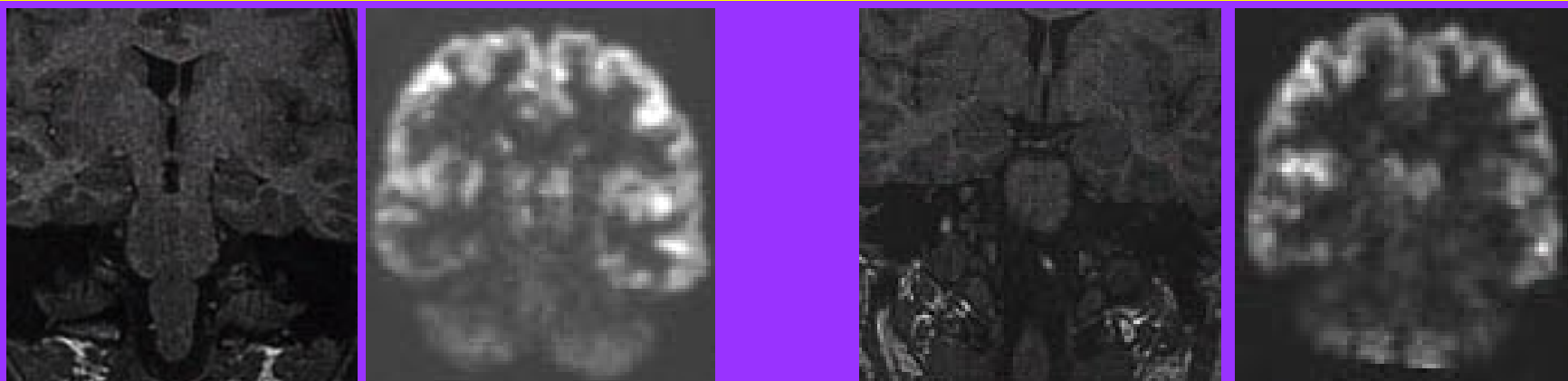


SIVD: Lateralization and Asymmetry

| SIVD | Concordant | Discordant | Non-lateralized |
|-------------------------------|-------------------|-------------------|------------------------|
| FDG-PET | | | |
| Assymetry Index > 3 | 20 | 0 | 3 |
| HV | | | |
| AI > 8 % | 15 | 0 | 8 |
| NAA/(Cho+Cr) | | | |
| AI > 12 % | 14 | 1 | 8 |
| NAA | | | |
| AI > 12 % | 15 | 1 | 7 |

Sharma et al(2003)Slovenia Medica Informatica(in Press)

MRSI: Coronal T1 weighted images with MRS spectra (SIVD)



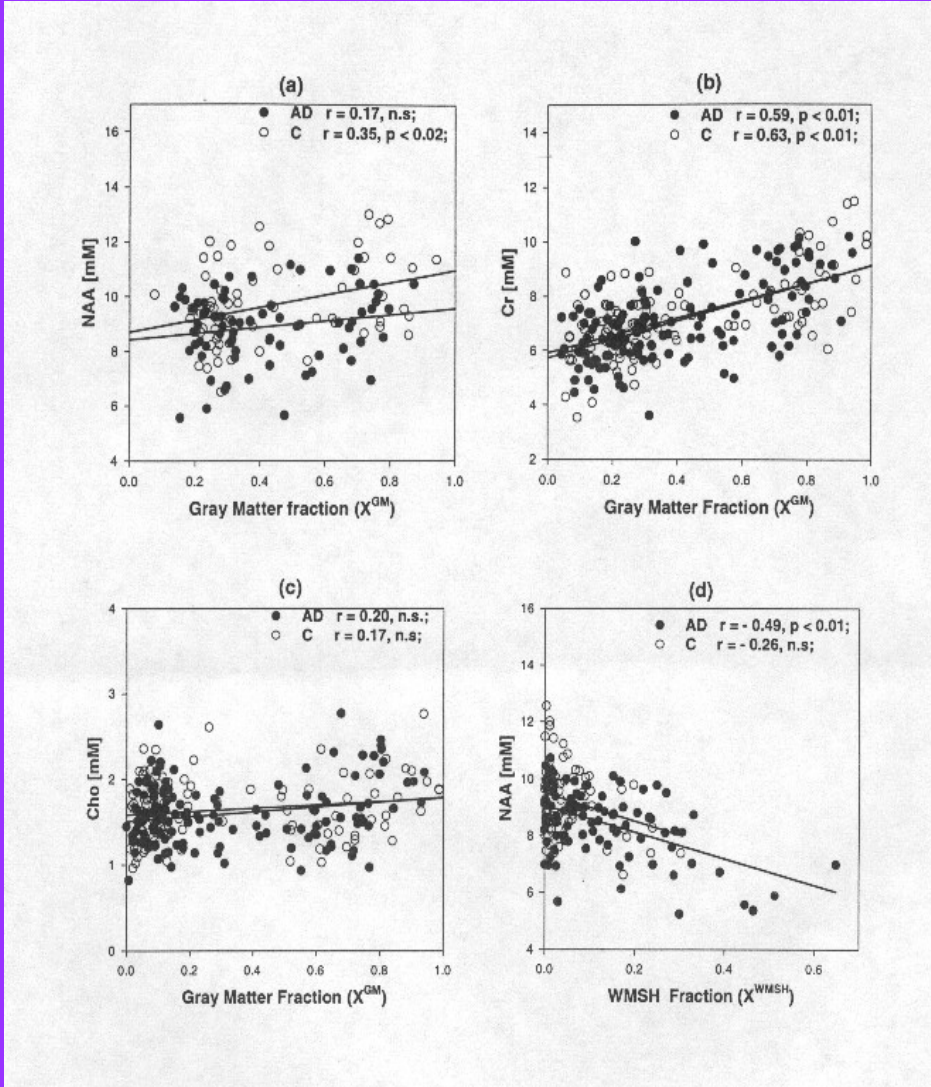
• Ipsilateral and contralateral differences in Choline peak

Regression Against Tissue Content and Histogram Analysis in AD

- **Regression analysis of $\text{NAA}^{\text{corrected}}$ as function of GM tissue fraction in each MRSI voxel.**
- **Histogram analysis of atrophy corrected NAA in parietal lobes (> 70 % GM) in AD**

Regression Analysis of Tissue Composition and Metabolites in AD

Nature Precedings : doi:10.1038/npre.2010.4317.1 : Posted 28 Mar 2010



Conclusion

- **Metabolites measure optimum lateralization**
- **Decreased NAA for hippocampus atrophy**
- **Bilateral abnormalities by Asymmetry Index**
- **Lateralization and discordance of lobes:**
Atrophy; HV; T2; NAA; NAA/(Cho+Cr)
- **Multi-slice MRI approach for Asymmetry Index**
- **MRI-defined two AD and SIVD disorders**

Where we go from here?

Specific Aim 1:

Better spectral MRSI resolution

Specific Aim 2:

Amyloid proteins and CSF proteins

Specific Aim 3:

Dementia neuropsychological classification and metabolite regional differences and Source of NAA

Specific Aim 4:

**Gene expression and regional metabolite
NAA/Cr+Cho ratio**

Technique Development for Spectroscopic Imaging Display(SID)

- SNR enhancement, B_0 inhomogeneity, 1st/2nd order phase correction, minimization of operator interaction
- Parametric Automated spectral simulation
- *a priori* information of metabolites
- voxel-by-voxel training data for tissue composition(CSF nulling)
- Point Spread Function and peak-overlaps
- Data Processing in the X, Y co-ordinates
- Corregistration and segmentation
- Correction of lineshape variation due to T_2^*

MRI + MRSI: Epilepsy

- **Neocortical Epilepsy**
- **Mesial Temporal Lobe Epilepsy**
- **Post-operative Mesial Lobe Epilepsy**

MRI: DSE and MP-RAGE images

- Oblique axial **Double Spin Echo(DSE)** T2 weighted imaging at TR/TE1/TE2 300/20/80 ms; resolution 1 x 1.4 mm²; 48 slices.
- 3D T1 weighted **MP-RAGE** acquisition TR/TI/TE 10/250/4 ms; flip angle 15; resolution 1 x 1 mm²; 48 slices.

Combined (multisection FLASH + PRESS) MRSI in mTLE

- Unilateral mTLE (ipsilateral side)
- **PRESS** volume pre-selection on hippocampus (TR/TE=1800/140 ms; voxel 9 x 9 x 15 mm³); circular K-space encoding of 24 points
- **Multisection FLASH** (TR/TE=1800/140 ms; voxel 8 x 8 x 15 mm³); circular K-space encoding of 36 points

Epilepsy may be better classified by H-1 MRI + MRSI

- **Neocortical Epilepsy**
- **Mesial Temporal Lobe Epilepsy(mTLE)**
- **Temporal Lobe Epilepsy(TLE)**

by

- **Ipsi- and contralateral localized changes**
- **Hippocampal Voluming**
- **PET and ectal EEG as clinical correlates of MRI and MRSI**

Hippocampus Lateralization by MRSI

- **NAA, Choline, Creatine and their ratio: NAA/Cr, NAA/Cho, NAA/(Cho+Cr) in left vs right hippocampus lobes in Epilepsy**

$$\frac{(M_{\text{contra}} - M_{\text{ipsi}})}{(M_{\text{contra}} + M_{\text{ipsi}})} \times 100 (\text{patients})$$

- **Asymmetry Index = -----**

$$\frac{(M_{\text{left}} - M_{\text{right}})}{(M_{\text{left}} + M_{\text{right}})} \times 100 (\text{controls})$$

M is metabolite concentration

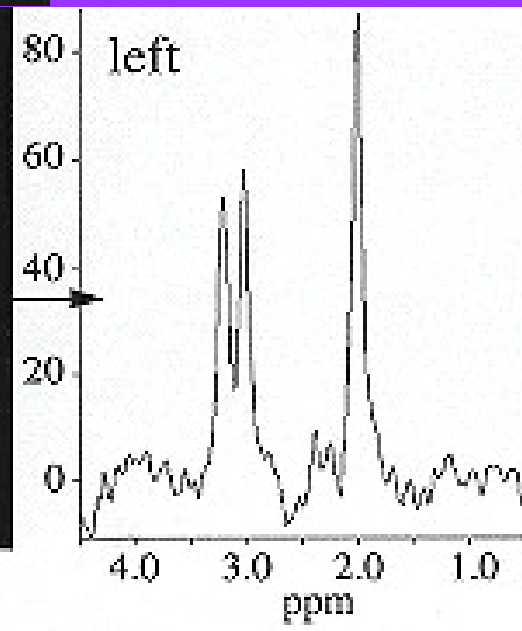
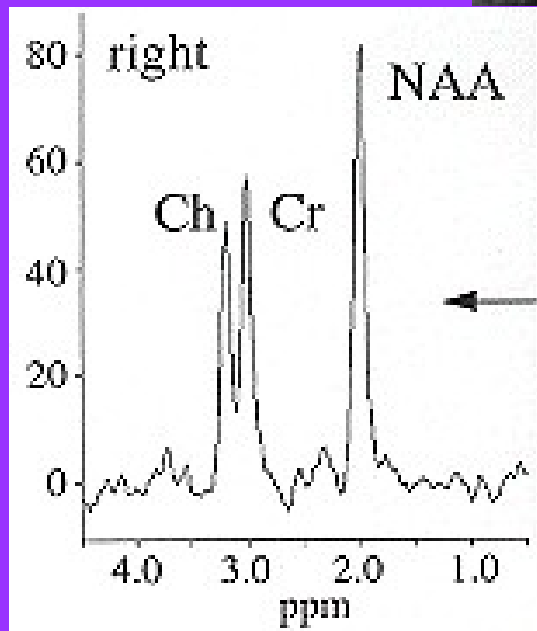
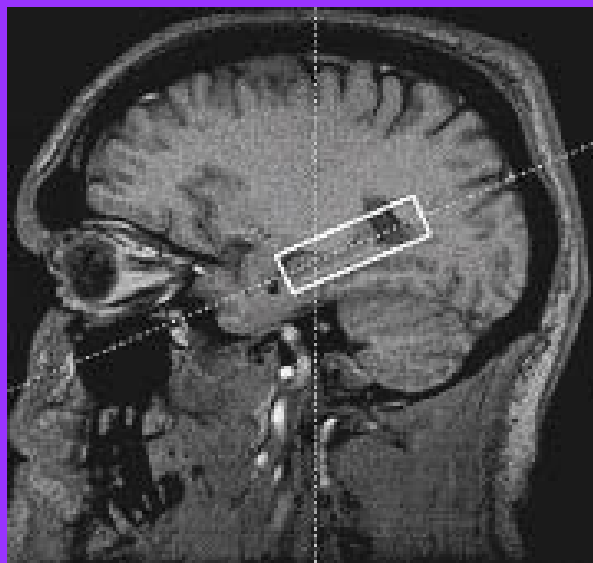
Criteria for Post-operative mTLE evaluation by H-1 MRI + MRSI:

- Percent Hippocampal volume reduction
- Reduced NAA/Cr
- Reduced NAA/Cho
- % Gray matter + % White matter

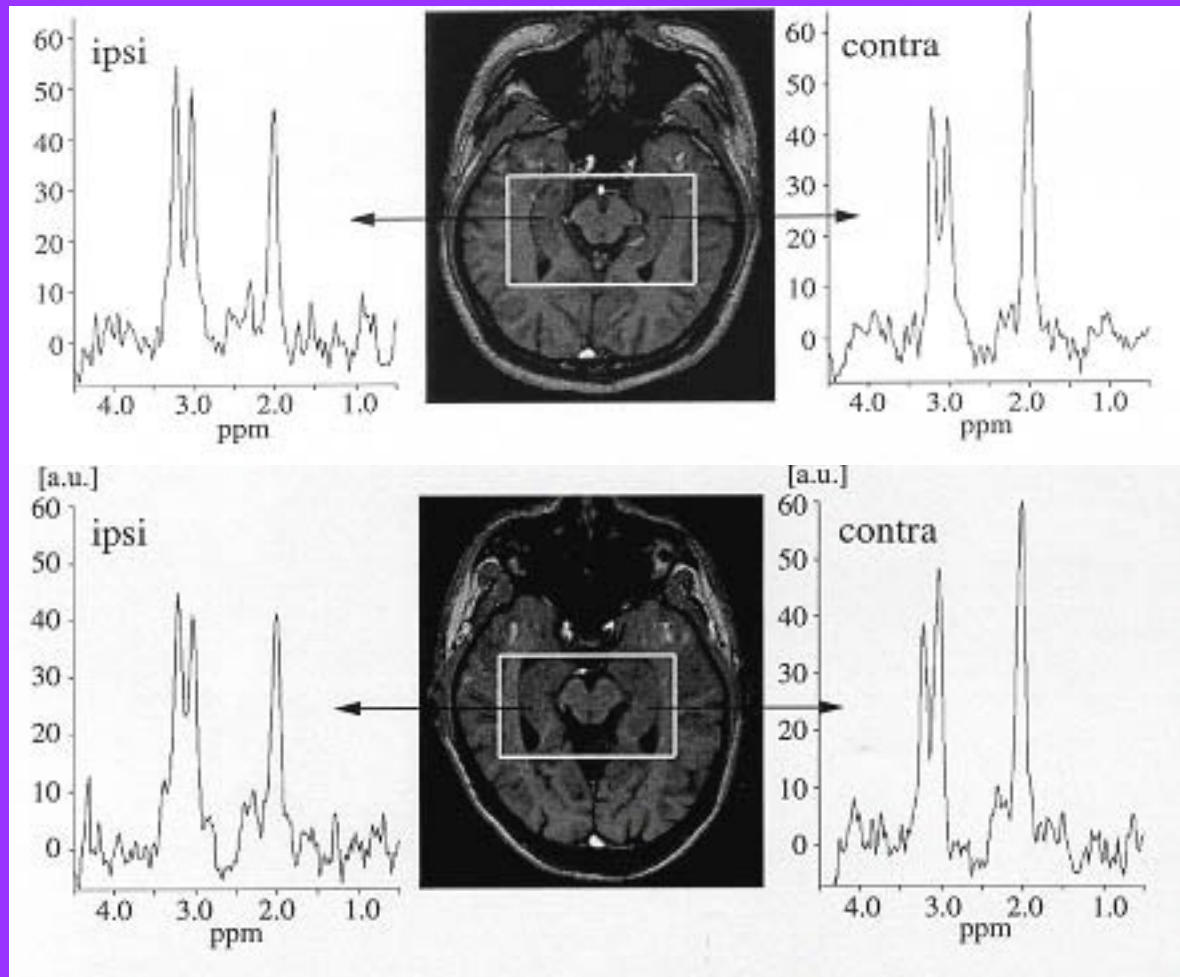
Characteristic changes in:

- Temporal lobectomy - Class I
- Partial seizures - Class II
- Complex partial seizure - Class III

PRESS: H-1 MRS of Hippocampus

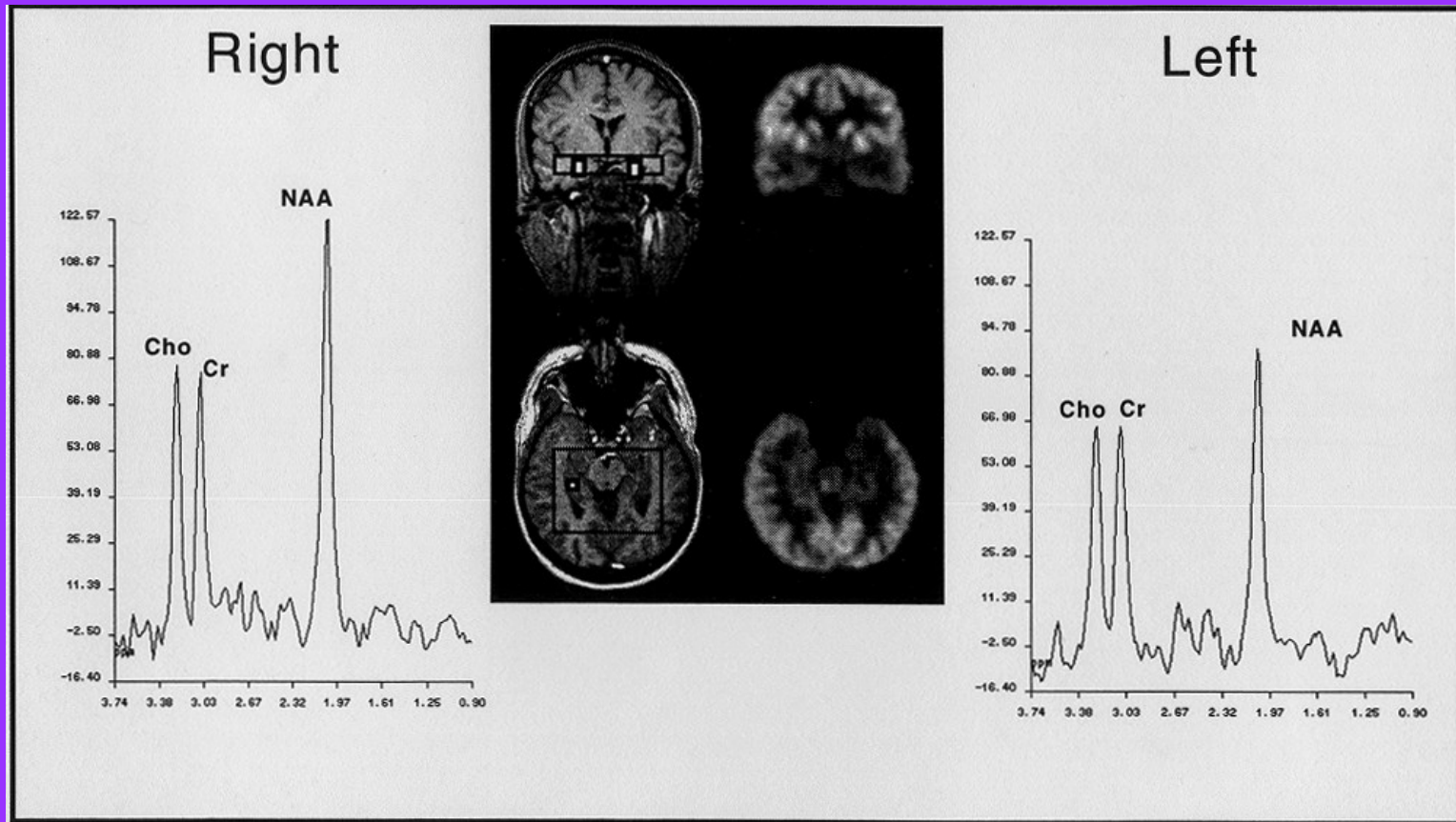


Transverse FLASH(Fast Low Angle Shot): TLE ipsilateral vs. contralateral metabolites



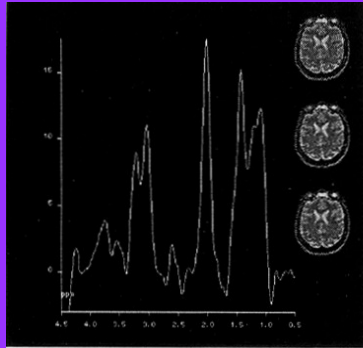
PRESS: Reduced NAA and Increased Cho in contralateral side

Nature Precedings : doi:10.1038/npre.2010.4317.1 : Posted 28 Mar 2010

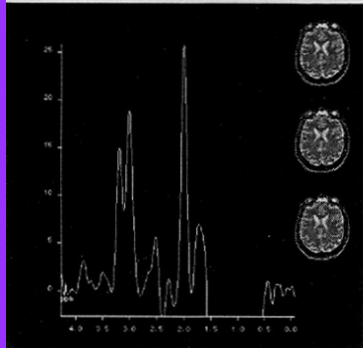


Sharma et al (2002) Slovenia Medical Informatica(in press)

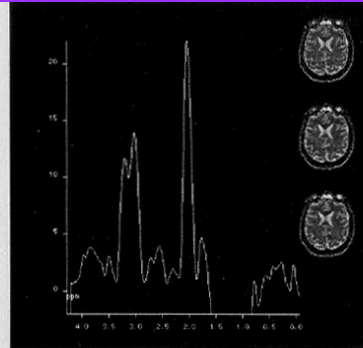
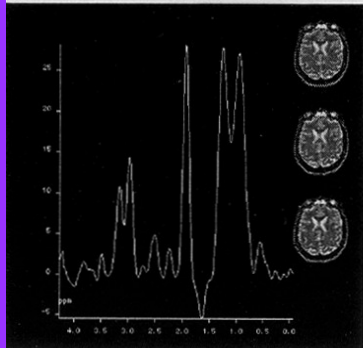
Voxelwise metabolite spectral peaks



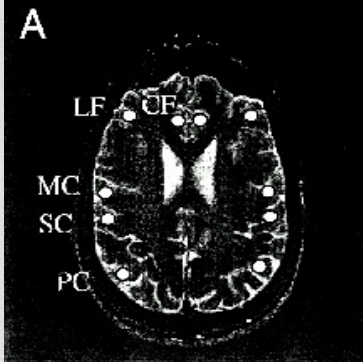
V1r



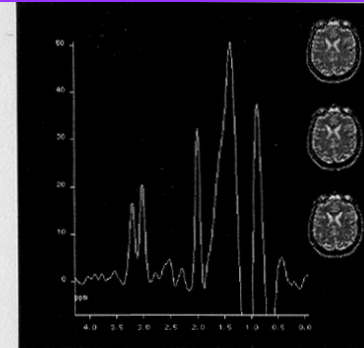
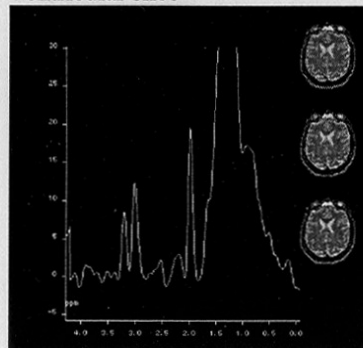
V2r



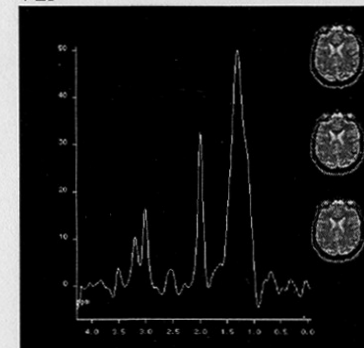
V1l



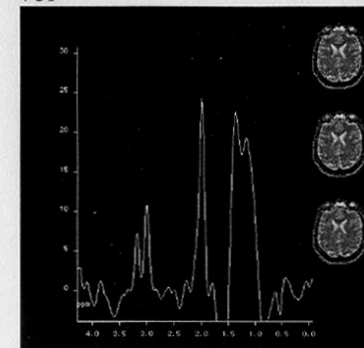
ventricular slice



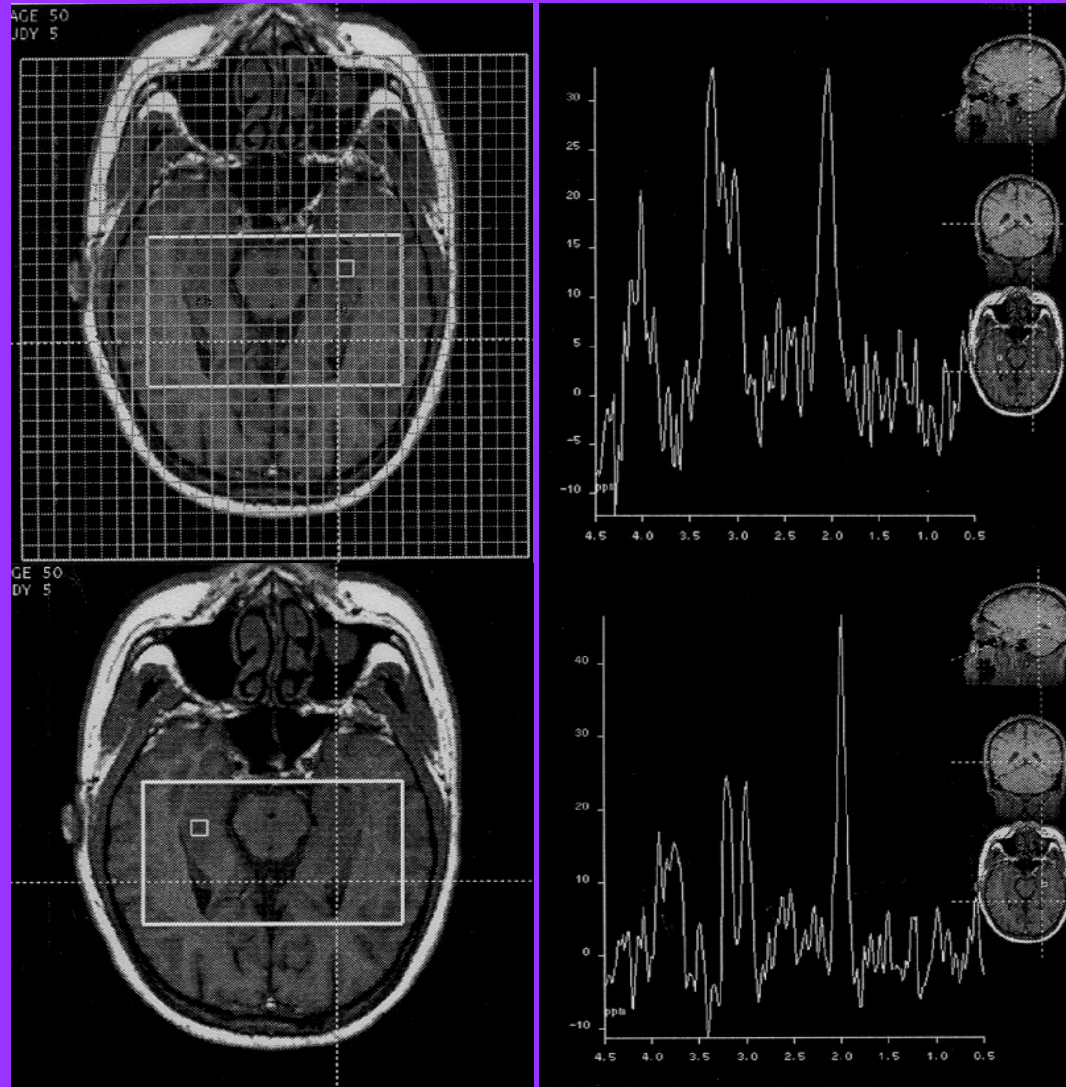
V2l



V3l

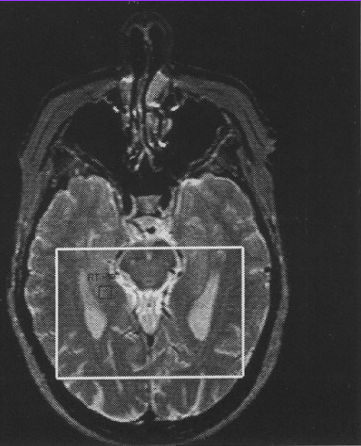


CSI: Metabolites in hippocampus left vs right lobes

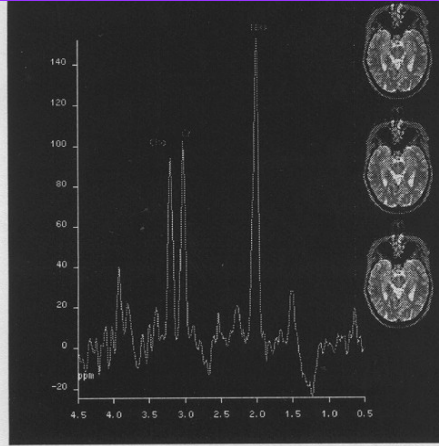


Hippocampus Volume Reduction and PRESS-MRS Lipids and Choline

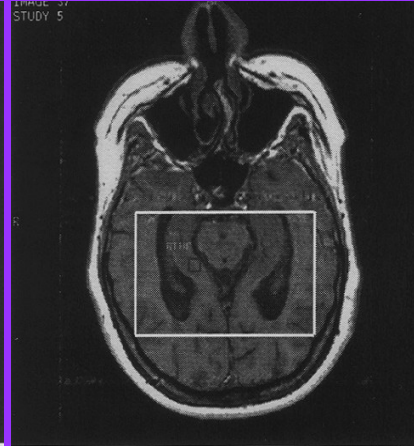
Nature Precedings : doi:10.1038/npre.2010.4317.1 : Posted 28 Mar 2010



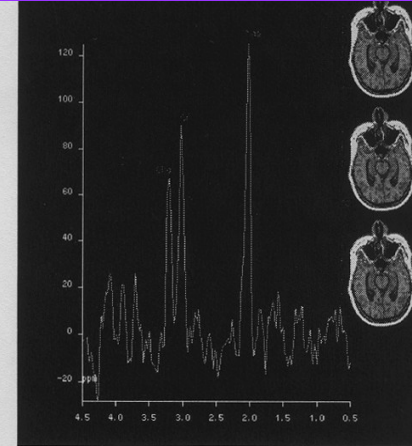
(A)



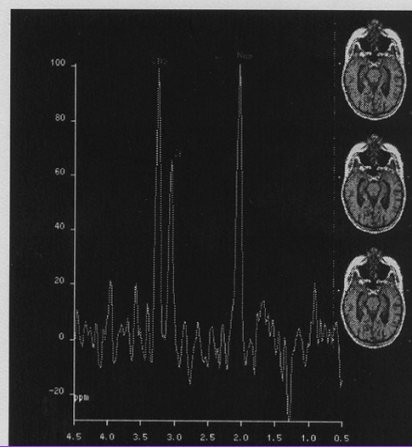
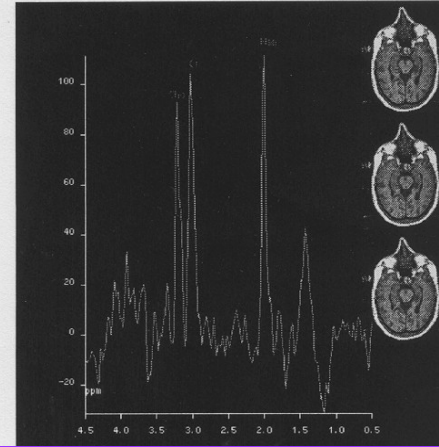
(B)



(A)



(B)



MRSI: Neuro-metabolites in Epilepsy

| Subjects | | NAA(mM) | NAA/(Cr+Cho) |
|----------------|---------------|------------|--------------|
| Controls(n=16) | | 11.6 ± 1.3 | 0.82 ± 0.06 |
| NE(n=8/10) | Ipsilateral | 12.3 ± 1.9 | 0.79 ± 0.1 |
| | Contralateral | 11.4 ± 2.7 | 0.78 ± 0.1 |
| mTLE(n=23) | Ipsilateral | 8.5 ± 1.3* | 0.62 ± 0.1* |
| | Contralateral | 9.6 ± 1.3* | 0.72 ± 0.1* |

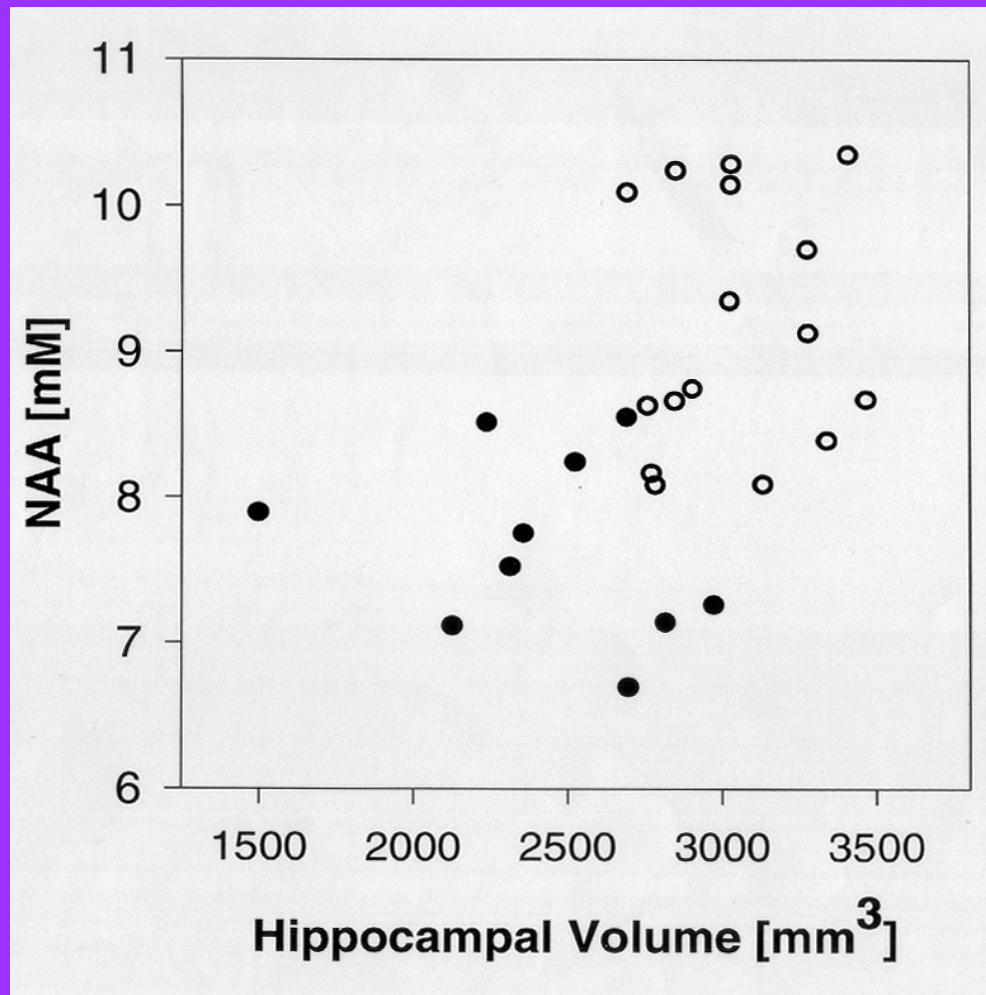
Ipsi- and contralateral NAA concentrations and NAA/(Cr+Cho) ratios(mean ± SD) in the hippocampus of NE and mTLE patients in comparison with controls (p values comparison in patients with controls). *indicates p value 0.001.

MRSI and Asymmetry Scores in Epilepsy

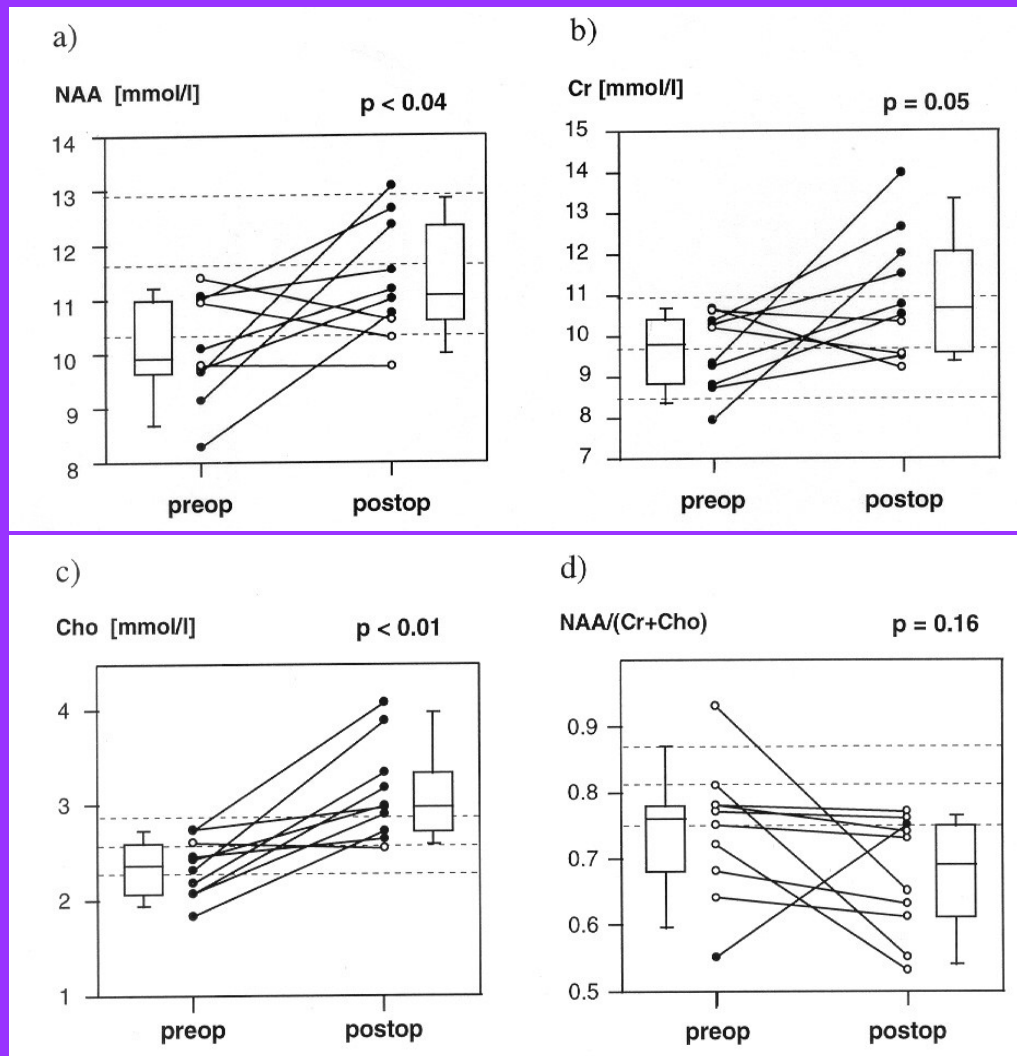
| Subjects | Asymmetry (NAA) | Asymmetry NAA/(Cr+Cho) | absolute | absolute |
|----------|----------------------------|---------------------------|--------------------------|---------------------------|
| | | | asymmetry NAA | asymmetry NAA/(Cr+Cho) |
| Controls | 4.85 ± 0.2 | 6.25 ± 0.63 | 5.65 ± 3.15 | 4.75 ± 3.95 |
| NE | 7.8 ± 4.6 | 8.01 ± 1.2 | 6.8 ± 5.71 | 6.75 ± 3.8 |
| | n.s. | n.s. | n.s. | n.s. |
| mTLE | 7.45 ± 2.27 (p < 0.001) | 7.4 ± 3.65 (p < 0.003) | 9.2 ± 6.4 (p < 0.001) | 9.07 ± 6.7 (p < 0.04) |

- **NAA and NAA/(Cr+Cho) Asymmetry Indices (mean ± SD) of NE and mTLE patients in comparison with controls (p values compared with controls).**
- **Absolute Asymmetry was independent of seizure focus**

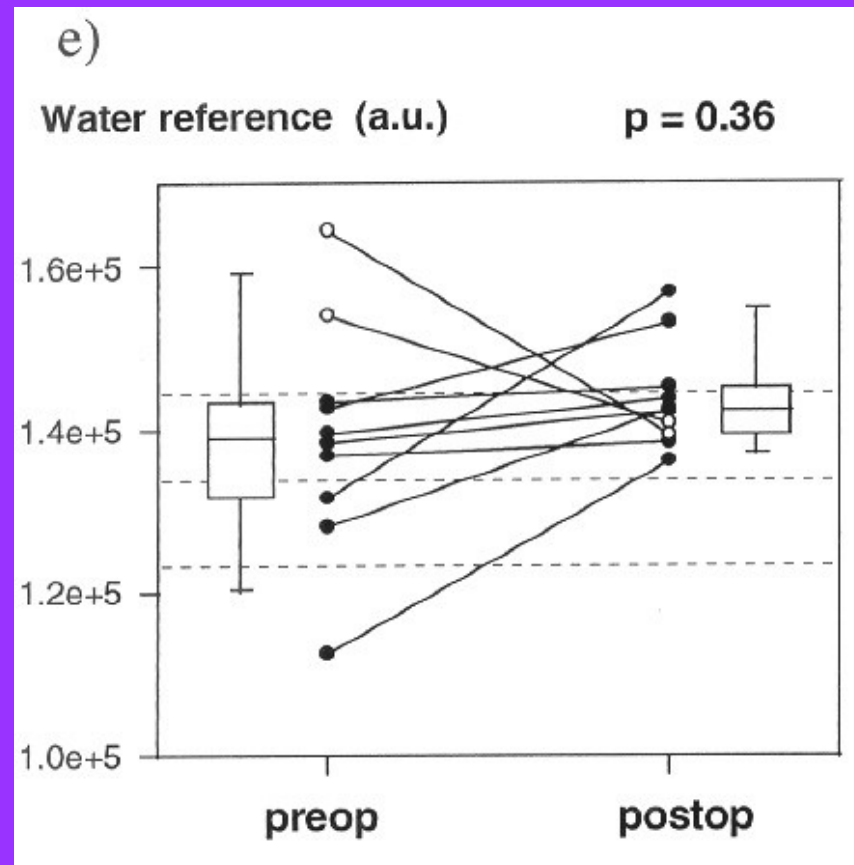
MRSI: NAA and Hippocampus Volume in Epilepsy



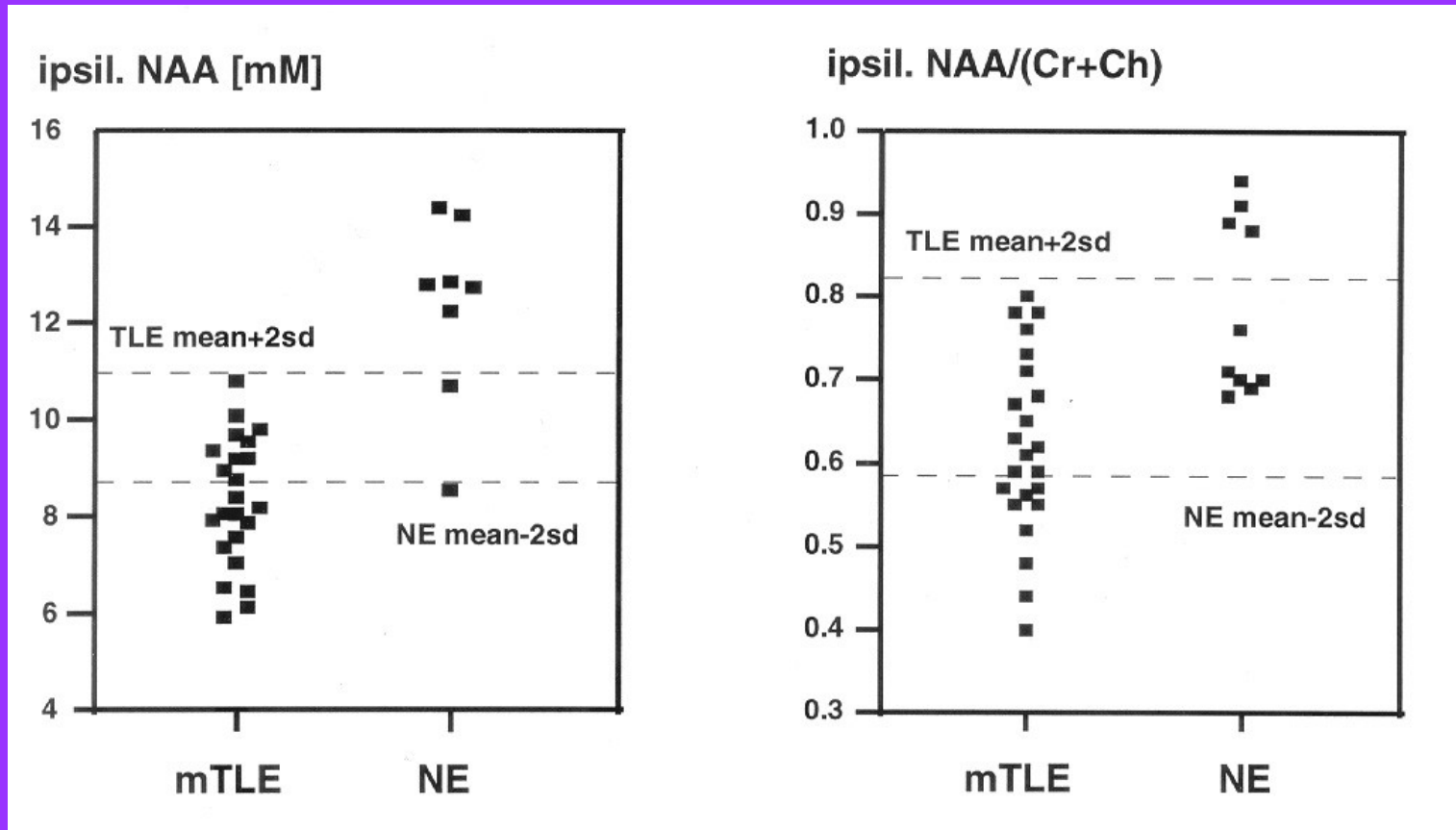
MRSI: Metabolites in Pre-op vs post-op Epilepsy



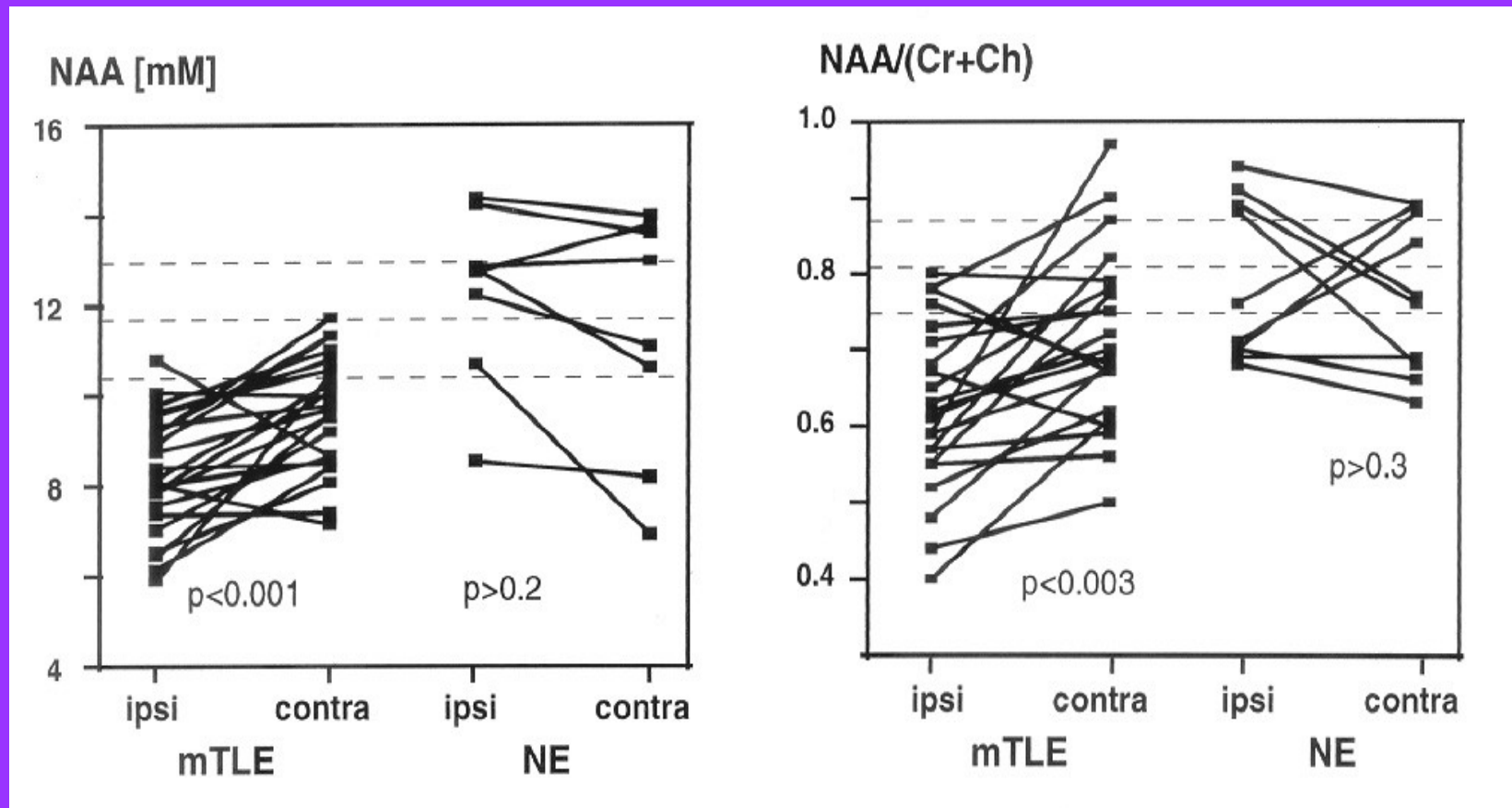
MRSI: Water as internal reference for metabolites in Epilepsy



MRSI: NAA and NAA/(Cr+Cho) in mTLE and NE Ipsilateral side

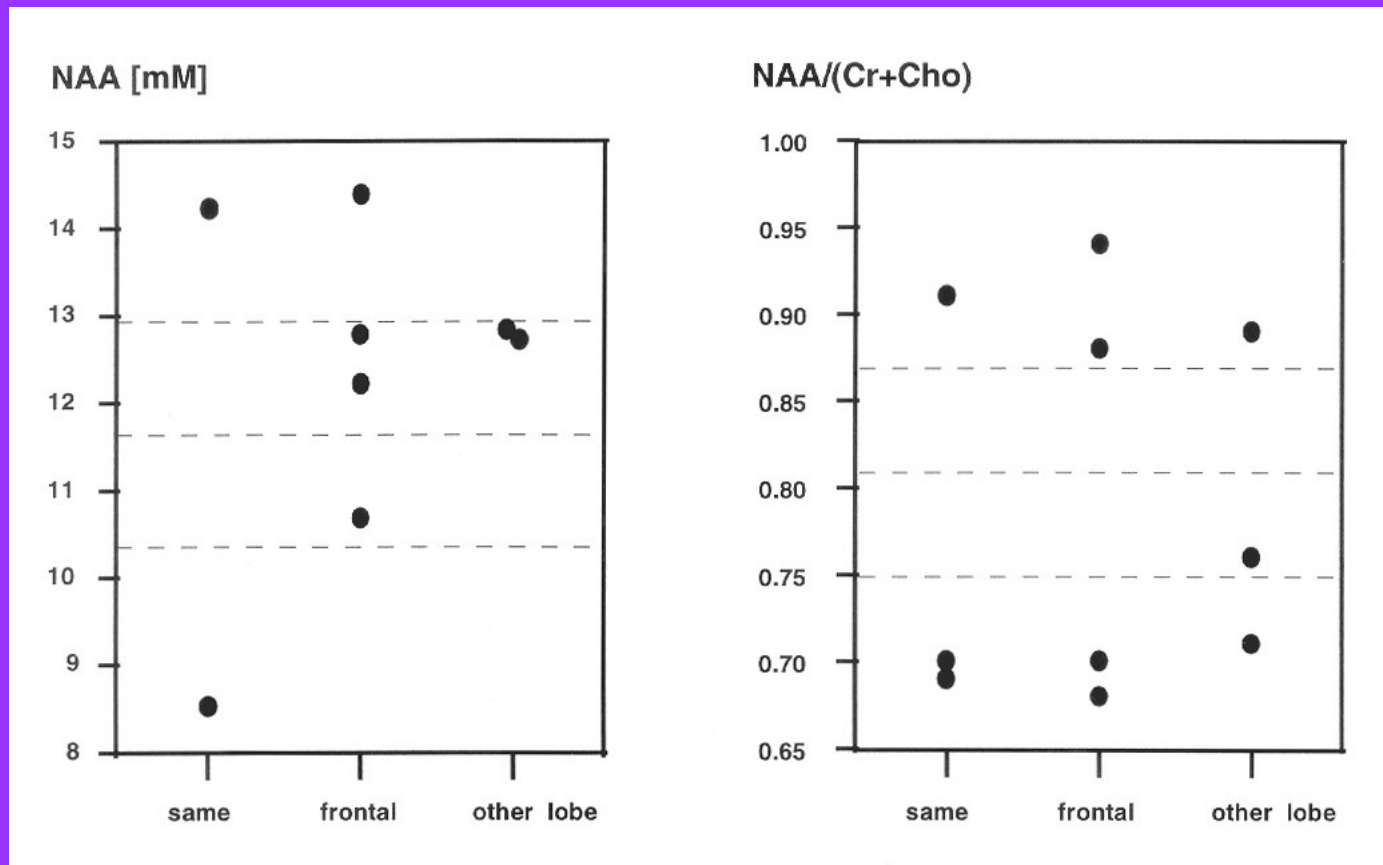


MRSI: NAA and NAA/(Cr+Cho) differ in NE and mTLE in ipsi- and contra-lateral sides

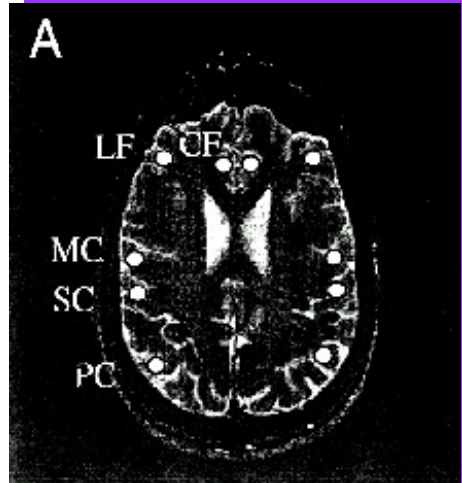
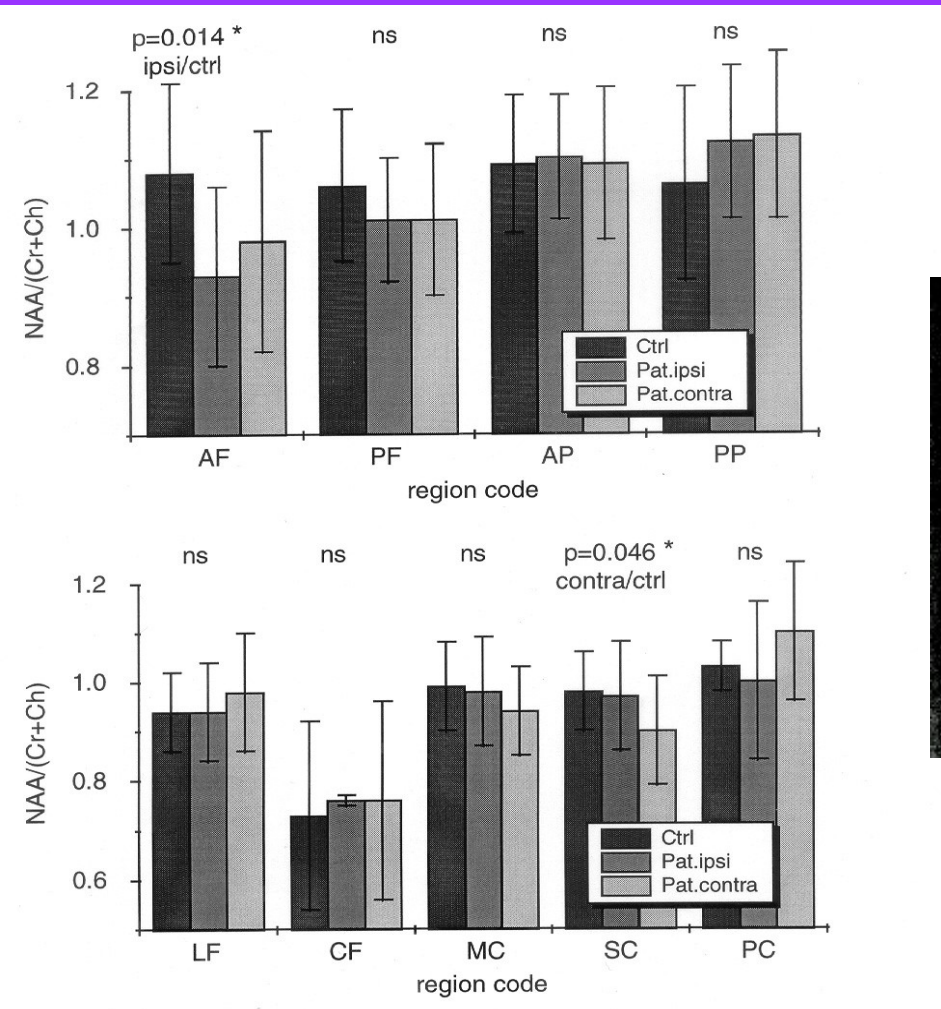


NAA, NAA/(Cr+Cho) in Frontal Lobe Epilepsy

Nature Precedings : doi:10.1038/npre.2010.4317.1 : Posted 28 Mar 2010



NAA/(Cr+Cho) in Control vs Epilepsy ipsi- and contra-lateral sides at different brain regions



Conclusion

- **Metabolites measure optimum lateralization**
- **Decreased NAA for hippocampus atrophy**
- **Bilateral abnormalities by Asymmetry Index**
- **Relationship of MRS peaks with post-op surgery seizure condition(bilateral abnormalities)**
- **Lateralization and discordance of lobes:
Atrophy; HV;T2; NAA; NAA/(Cho+Cr)**
- **Assessment of seizure focus spread**

From here where we go?

Specific Aim 1:

Absolute concentration of metabolites and 2D MR spectroscopy(COSY) information

Specific Aim 2:

Lateralization and discordance criteria

Specific Aim 3:

Sensitivity of the techniques by gradient correction, better Rf coil design

Specific Aim 4:

Alternative approach of FDG/PET and perfusion/diffusion weighted MRI

Specific Aim 5:

Robust software for registration-segmentation and hippocampus voluming

MRI and MRSI in Multiple Sclerosis

Introduction

- Multiple Sclerosis, a neurodegenerative disorder with relapsing remitting course is manifested as progressive **lesions** around ventricles.
 - i. Blood-brain barrier altered--> demyelination of axon-->MS lesion enhancement(MRI-defined Perivenous inflammation)
 - ii. Non-inflammatory, **lipid rich WM and GM(No MS lesion)**
- **Serial MRI/MRSI data offers:**
 1. MS lesion volumetry
 2. Brain GM and WM composition
 3. Comparison of H-1 MR metabolite images,
 4. *in vivo* H-1 MR spectral analysis in different regions of brain

Hypothesis

- **Mobile lipids release at the cost of NAA (without BBB breakdown) initially followed by amino acid release.**
- **Multiple Sclerosis may be non-inflammatory with lipid-rich Normal Appearing White Matter or Gray Matter (NAWM or NAGM)**

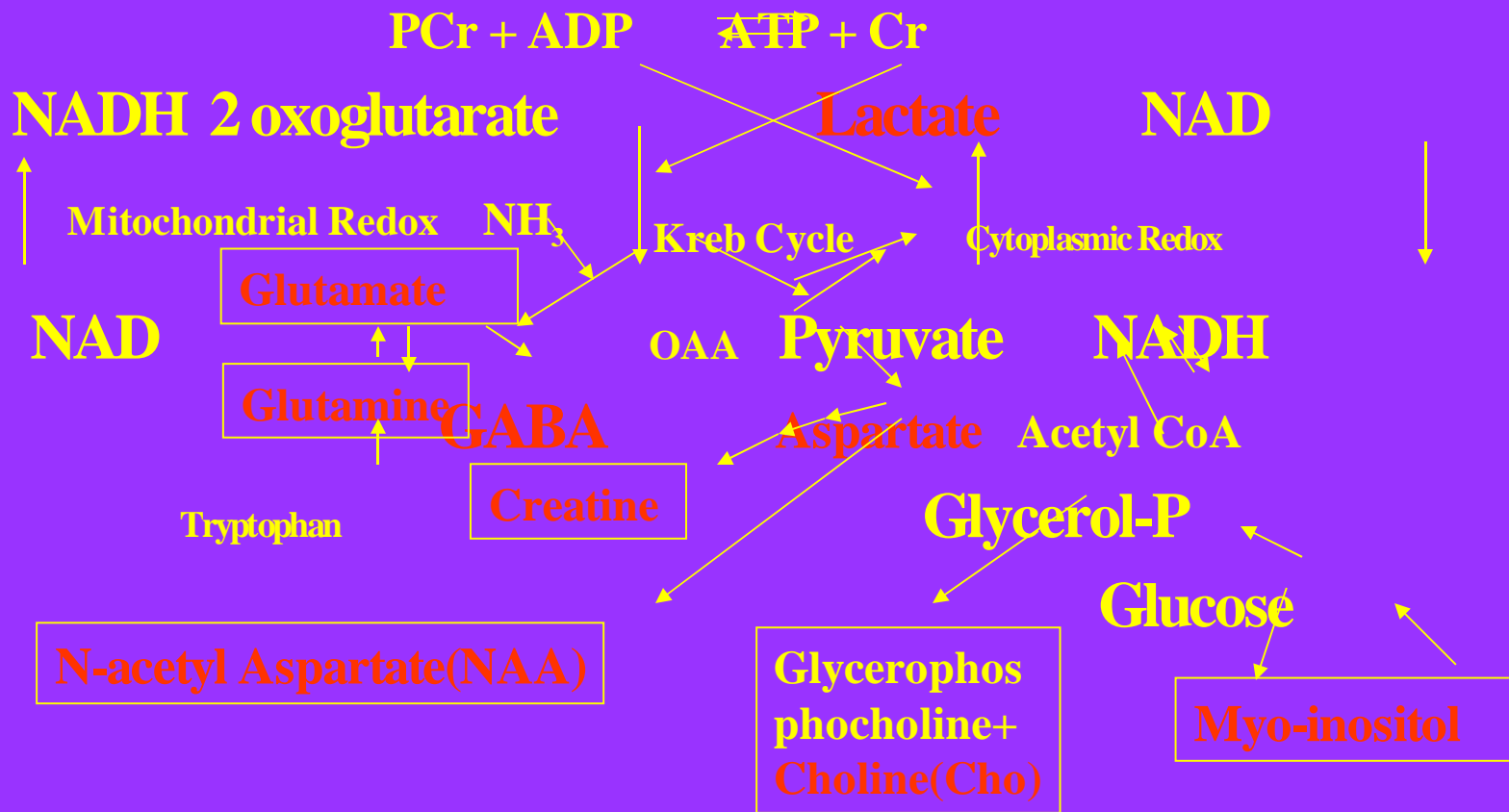
Approach of MRI/MRSI

- Brain and MS lesion tissue composition by **MRI** in serial scans:
 - Tissue segmentation:
 - Supervized and Automated Method:
 - MS lesion volumetry by ‘seed growing algorithm’
 - Brain WM/GM feature-space (Perzen Window) volumetric analysis
 - Gd-lesion enhancement by saturation bands & Gradient dephasing
 - Tissue 3D Registration:
 - Intrahemispheric Fissure(IHF) and Edge-detection method:
 - Search window
 - Rotations about z and y axis
 - Spatial Transformation and Image reconstruction
- Regional metabolite differences by **MRSI**:
 - Metabolite image generation at TE=20 ms
 - Standardization of MR peaks (*a priori* information based parametric fitting)
 - Automated spectral analysis of all image voxels of MRSI data set in iterative manner (nonparametric baseline and parametric fitting of model metabolites)

MRI+MRSI in Multiple sclerosis

- **Serial studies of multiple sclerosis(MS) lesion load(MRI)**
- **Serial brain metabolite imaging (Lipids)**
- **1.5 T MR chemical shift imaging(CSI)**
- **MRI and MRSI Co-analysis**

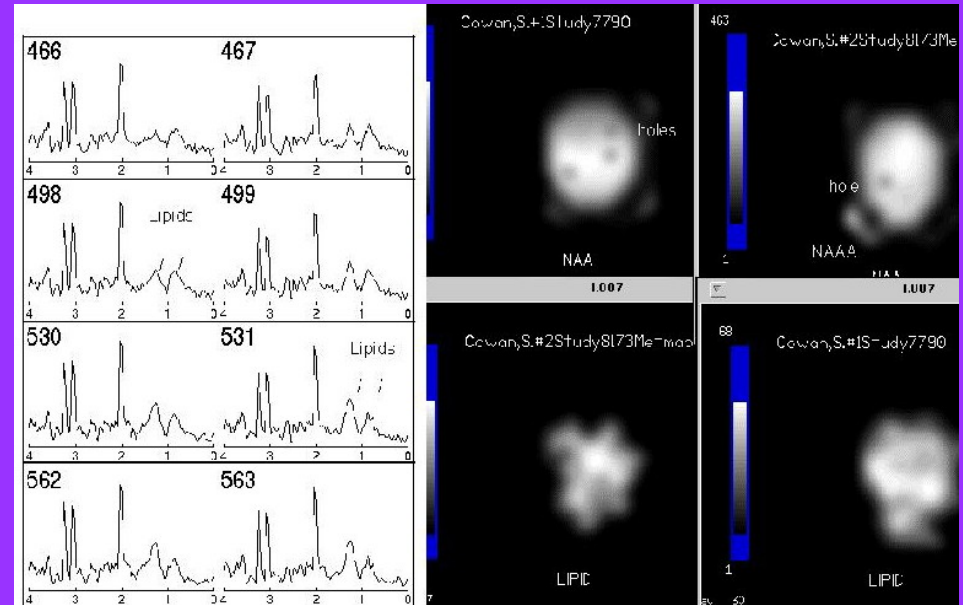
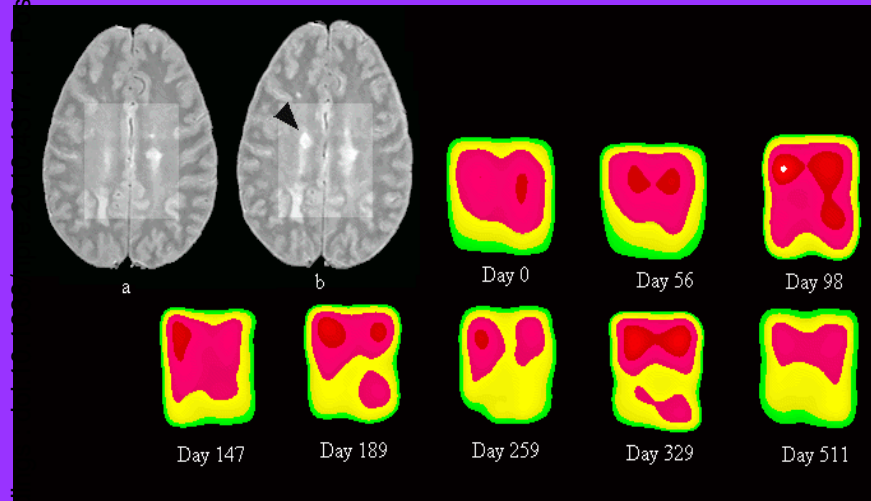
MR visible metabolites in brain



MRSI postprocessed metabolite NAA, Choline images in MS

Presented 28 Mar 2010

Nature Preced



- Enhanced Mobile Lipids (doublet peak) and Choline associate with Reduced NAA

MRSI: Lipids in serial MS lesions

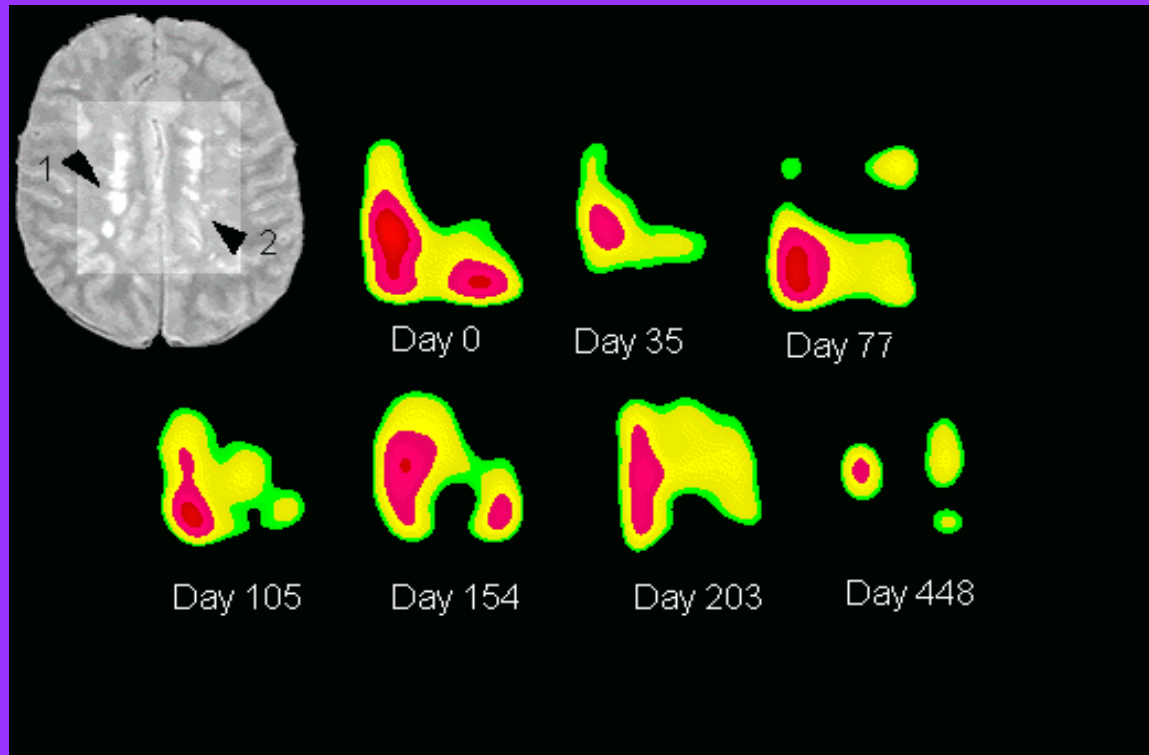
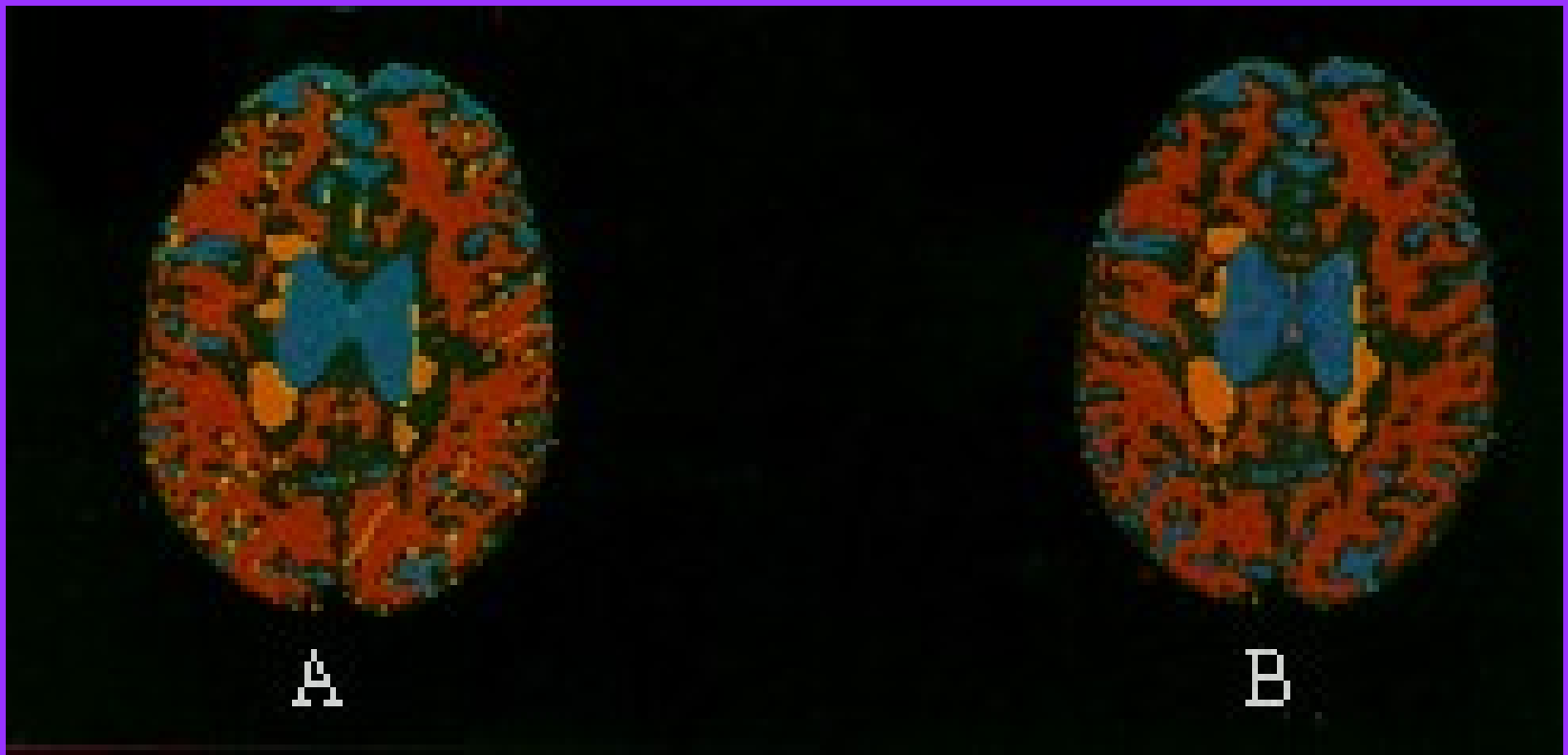


Figure 7: Spectroscopic VOI highlighted on 15 mm thick collapsed(left) slice image with arrowed enhancing lesions. Lipid distribution in lesions shown in time dependent manner. Note lipid images in NAWM(day 0) and later enhancing lesions(after 35 days) and recovered later.

MRI+MRSI in MS

- **RF inhomogeneity correction**
- **3D image registration**
- **Metabolite concentrations and % tissue composition**
- **Automated segmentation of Gd-enhanced MS lesions**
- **Serial lipids and other metabolite changes in MS lesions**
- **Serial lesion volumetry**
- **GM seated MS lesions and NAGM vs NAWM**

Step 1: Segmentation- Feature maps of different regions in MS brain

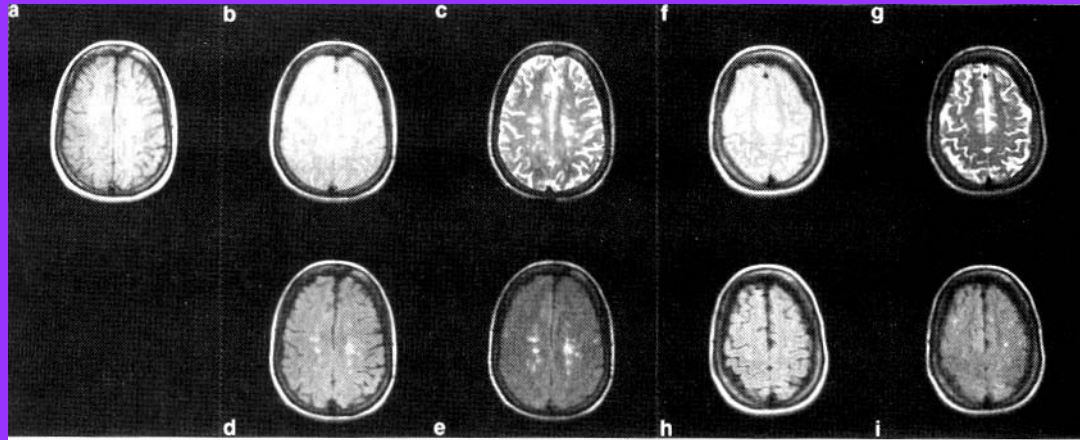


Automated segmentation of Gadolinium enhanced MS lesion



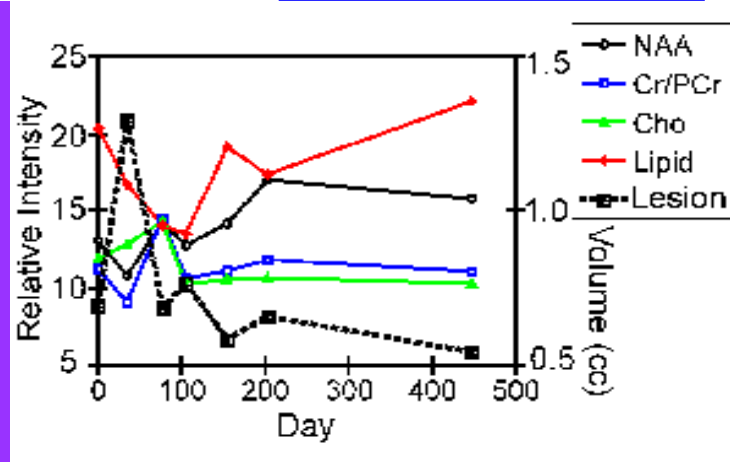
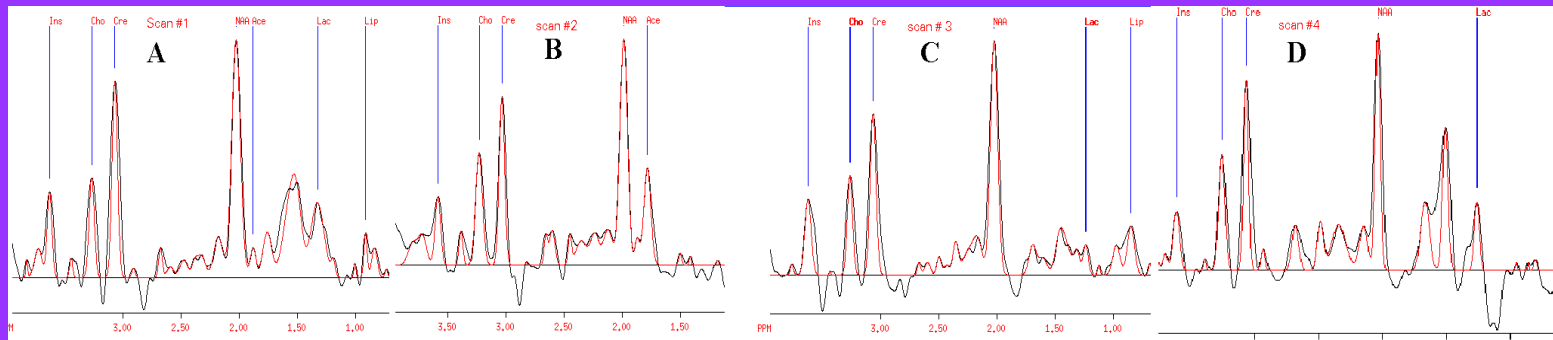
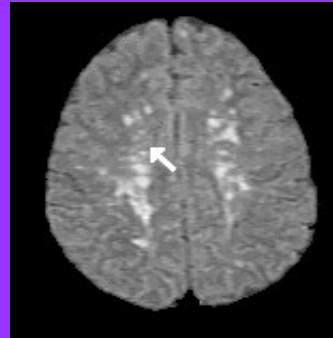
- **a. Raw Image; b. Removal of scalp; c. CSF suppressed; d. WM+GM+CSF suppressed; e. Partial Volume Average Corrected**

Step 2: Registration Technique

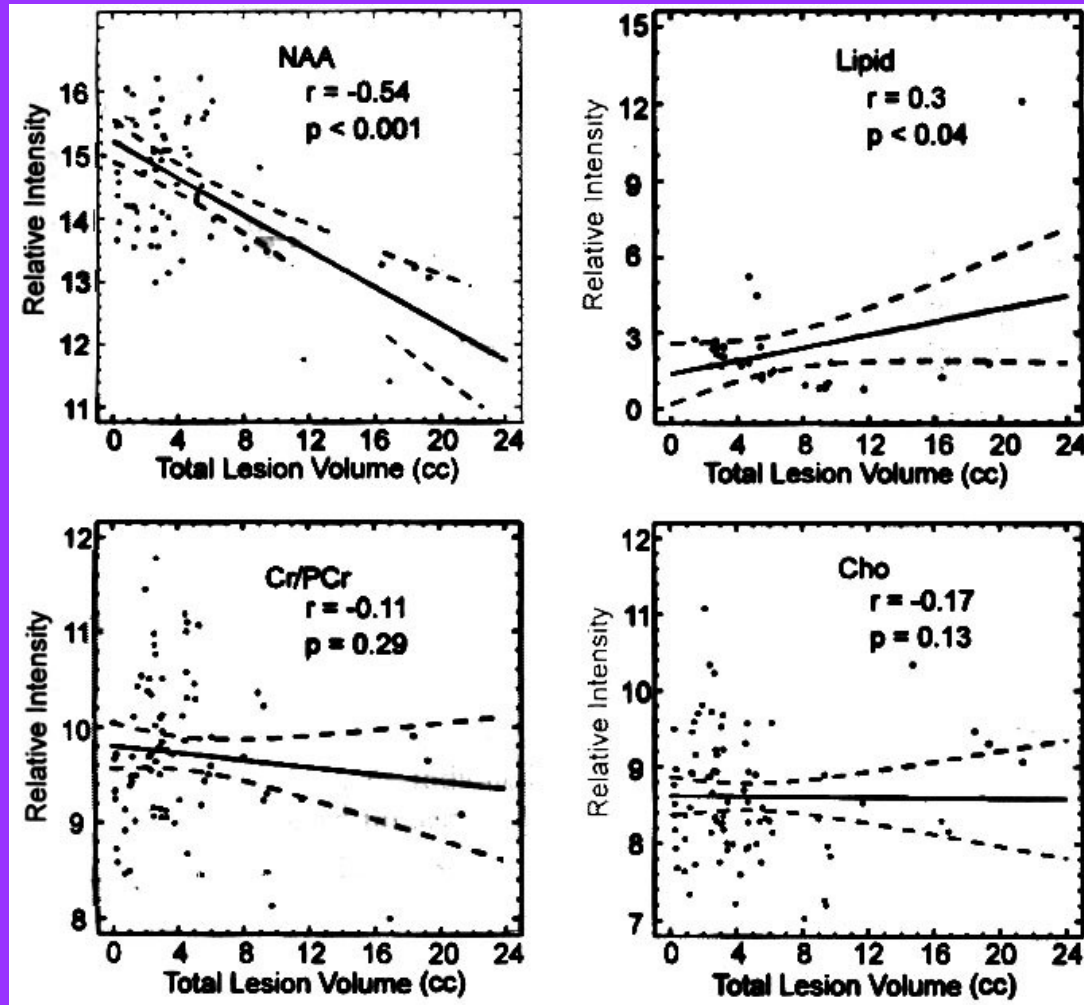


a. Post-contrast T1-w image; b-e. 4 FSE images with early and late echoes after registration; f-i. Same 4 images after registration

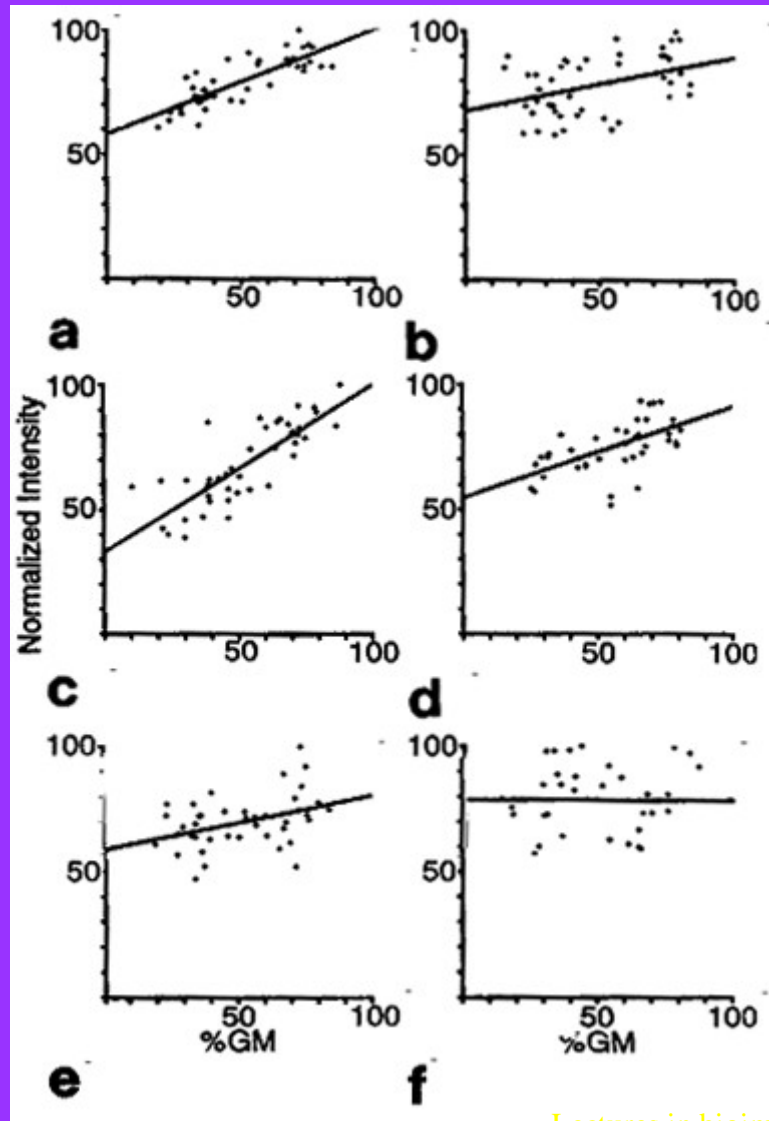
Time dependent metabolite changes in MS brain



Regression Analysis: Metabolites and MS lesion Volumes



Neurochemicals vs %GM in Normal Brain

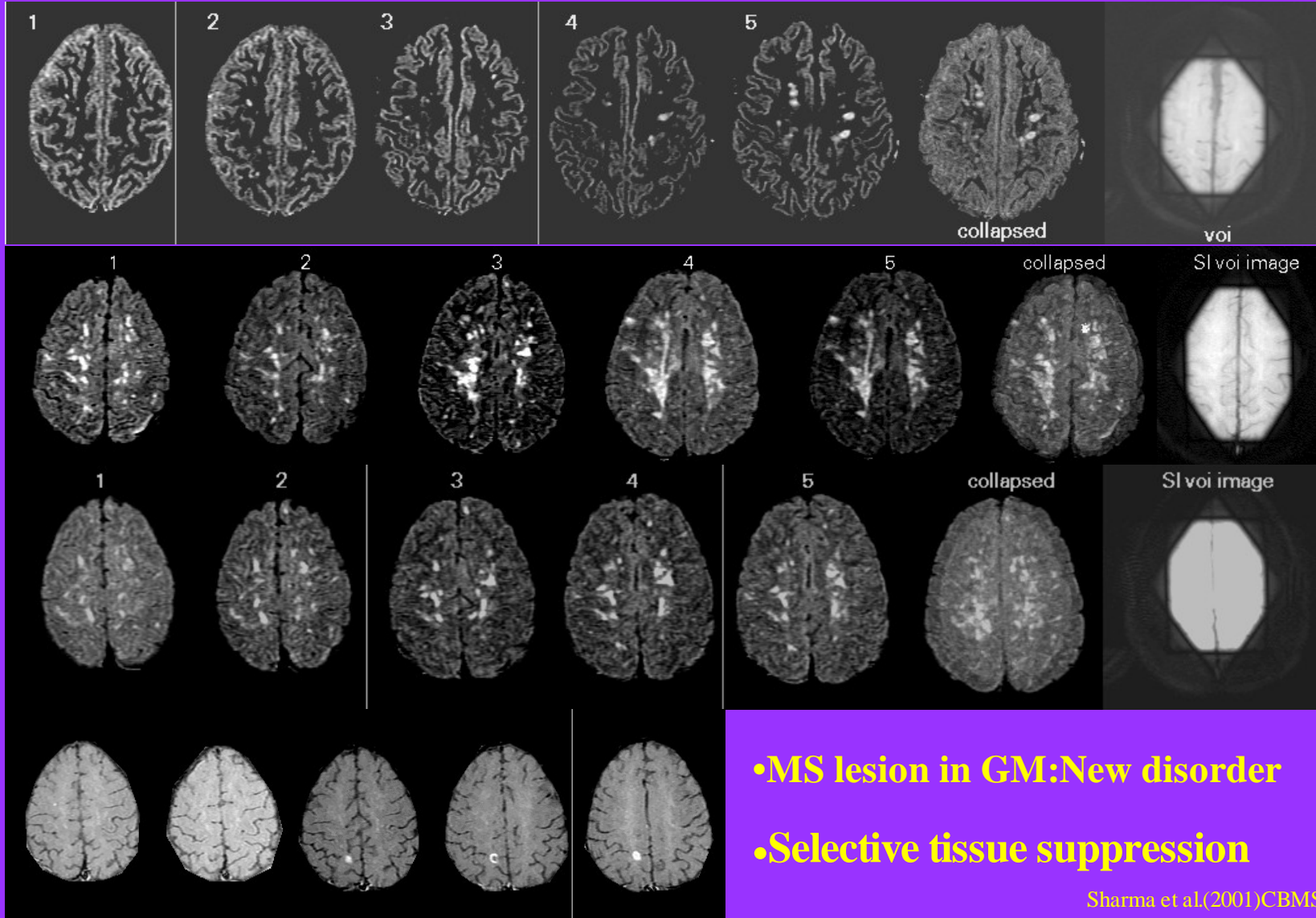


- Scatter plots of neurochemical vs % GM for:
- NAA:
 - a.($R=0.82, P < 0.001$);
 - b.($R=0.42, P < 0.003$)
- Cr/PCr:
 - c.($R=0.81, P < 0.01$);
 - d.($R=0.56, P=0.001$)
- Cho:
 - e.($R= 0.4, P < 0.009$);
 - f.($R=-0.01, P = 0.994$)

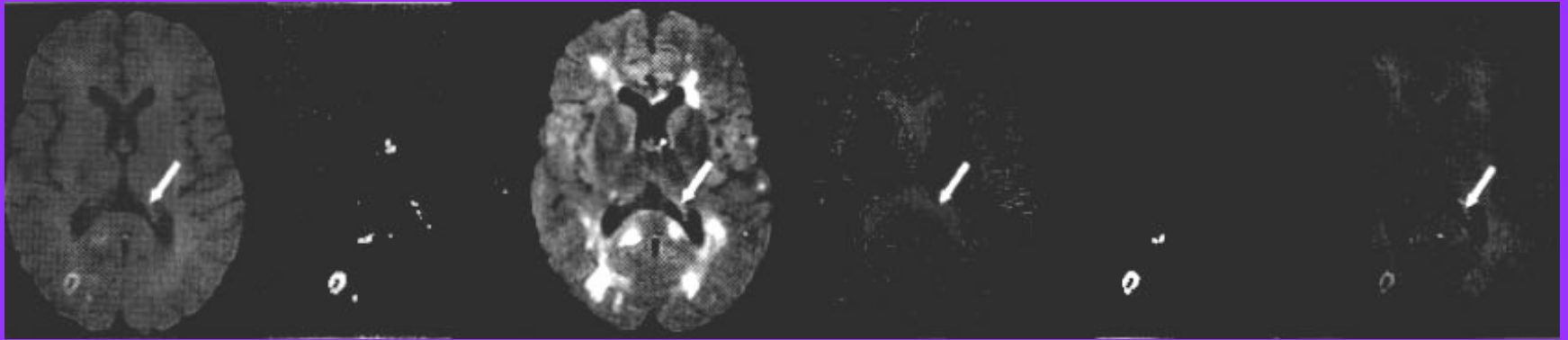
MS: Non-Inflammatory GM Lipid Disorder

- Multiple Sclerosis(MS) is known as demyelinating White Matter disorder
- Histology data suggests abnormal GM in MS
- Occasional presence of GM lesions on MRI

MS Lesion Showing Abnormal Metabolites in Pre- and Post-contrast GM and NAGM



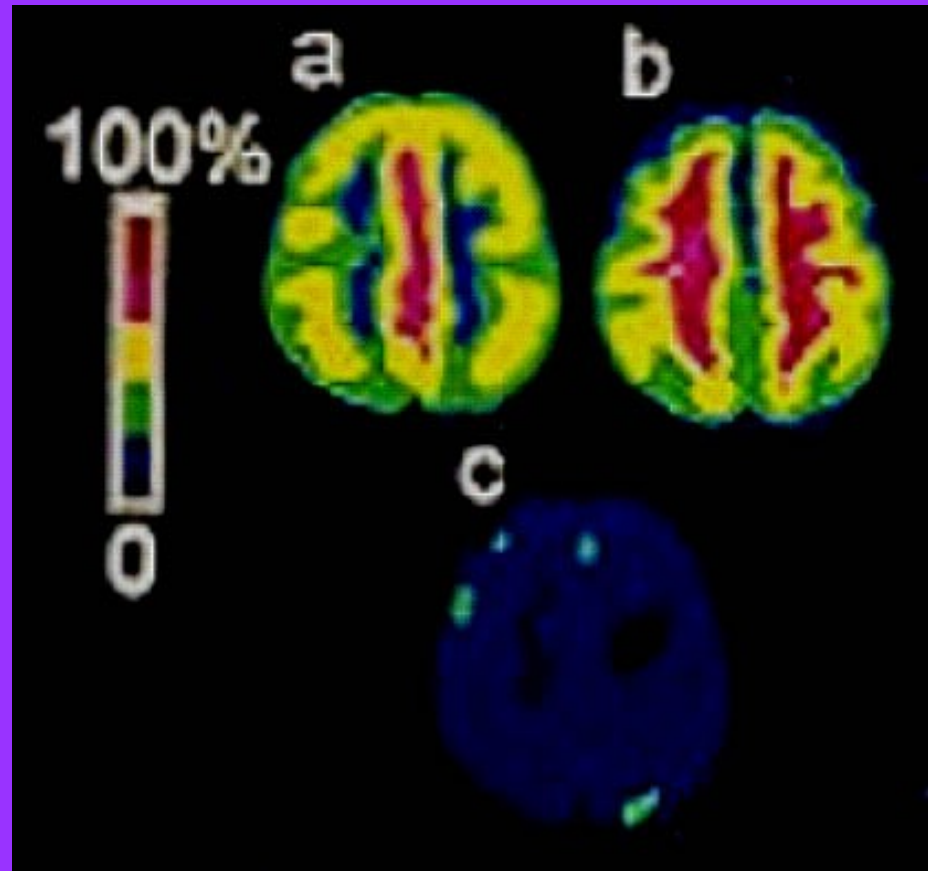
Gd-Contrast enhancement on MR slice showing GM Lesion



- GM+WM+CSF suppression visualize GM seated lesion after Post Gd- injection

Selective Tissue Segmentation Method

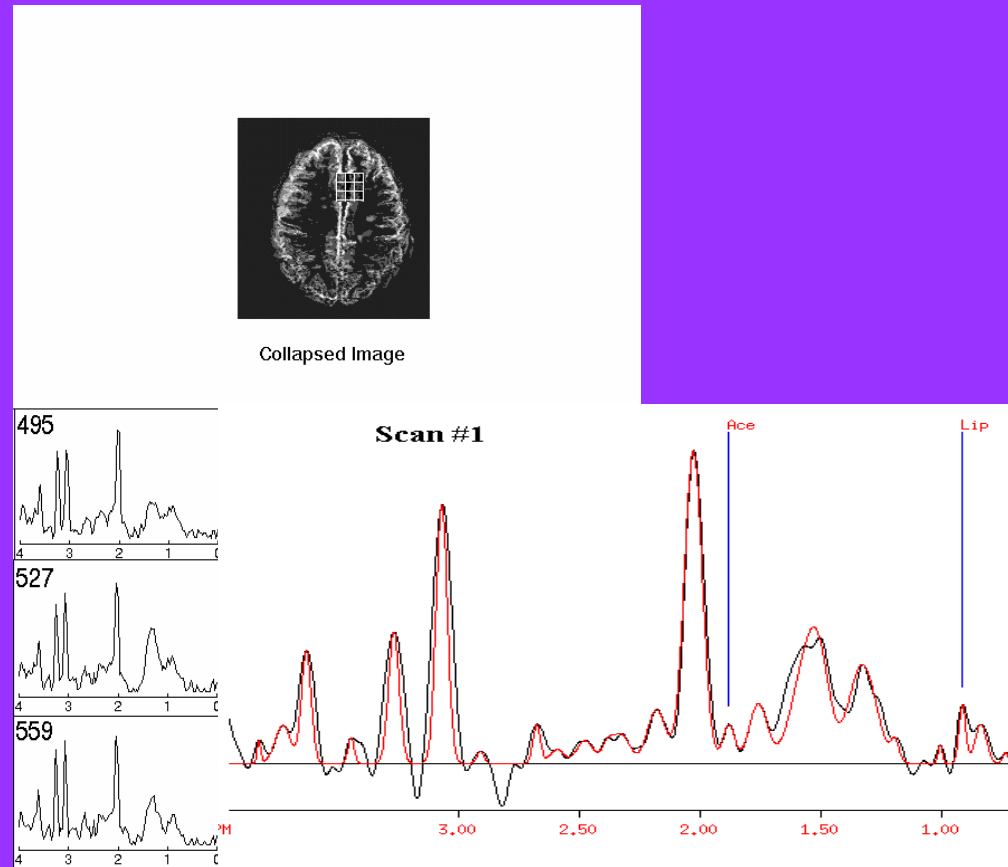
- Segmented images of GM(a); WM(b); CSF (c)
- 15 x 15 pixel² Gaussian-shaped filter with FWHM= 11 pixels applied



MS: Lipid and Lactate in NAGM

•MS Lesion not visible

•Lipid(doublet) and Lactate peaks



Distribution of lipid rich MS brain voxels in gray matter(GM) and normal Appearing Gray Matter(NAGM)

| Voxels in different regions in brain hemisphere | | | | |
|---|---------|----------|-----------------|-------------|
| | Frontal | Parietal | Fronto-parietal | Mid-fissure |
| Left GM | 16 | 5 | 6 | |
| NAGM | 29 | 14 | 7 | |
| Right GM | 18 | 5 | 8 | 210 |
| NAGM | 79 | 27 | 23 | |

Percent Tissue Composition of Brain GM and NAGM Regions in MS

| GM(n=65) | |
|----------|-------------------|
| %GM | 82.41 \pm 22.36 |
| %WM | 10.2 \pm 16.7 |

| NAGM(n=428) | |
|-------------|-------------------|
| %GM | 82.14 \pm 24.83 |
| %WM | 12.25 \pm 18.78 |

(Values shown as mean \pm 2 S.D. after CSF correction)

Absolute Concentrations of Metabolites in Spectroscopic Voxels in MS Patients

| MS Patients: | | |
|--------------|------------|-------------|
| | GM(n=65) | NAGM(n=428) |
| NAA | 10.02±3.95 | 10.8±4.6 |
| Cr | 8.39±3.16 | 6.73±1.36 |
| Choline | 6.43±2.96 | 3.76±1.19 |
| Lipids | 6.15±3.87 | 2.60±1.61 |

| Normal Volunteers:No lipids | | |
|-----------------------------|------------|-----------------|
| | GM(n=50) | All types(n=75) |
| NAA | 10.05±3.26 | 9.61±3.33 |
| Cr | 8.03±2.63 | 3.33±2.75 |
| Cho | 5.5±2.6 | 5.72±2.7 |

Outcome of MRI+MRSI Co-analysis

- **Quadruple Contrast images represent White matter and CSF suppression for clear MS lesion**
- **Cortical and subcortical lesions are normally MRI non-visible in GM**
- **MS lesion in GM represent strong peaks at 0.8-1.5 ppm(lipids and other metabolites)**
- **MS lesion rich GM represents 90 % GM and 10 % WM**
- **Neurochemical changes in GM and NAGM suggest membrane breakdown**

CONCLUSION

- **Gray Matter and Normal Appearing Gray Matter in MS are abnormal on MRS**
- **Neurochemicals in MS lesion-rich GM suggest MRSI measurement power for GM integrity**

Problem partly solved

- **Selective WM/GM suppression: (AFFIRMATIVE and QC pulse sequences)**

37,94,102(1997)

MRM

- **Lesion Volume, metabolite image-generation (APSIP, Image Viewer and MRIAP)**

MRM B, 106, 58(1995)

**AFFIRMATIVE: Attenuation of Fluid by East Inversion
Recovery with Magnetization Transfer Imaging with Variable
Echoes by FLAIR and MTC into spin echo**

QC: Quadruple contrast;

APSIP: Automated Spectroscopic Imaging Program;

MRIAP: MR Image Analysis Program

Where we go from here?

Specific Aim 1:

Better spectral MRSI resolution

Specific Aim 2:

Amyloid proteins and CSF proteins, Demyelination, inflammation and metabolite regional differences

Specific Aim 3:

Source of NAA, amino acids, Gene expression and regional metabolite NAA/Cr+Cho ratio

Specific Aim 4:

MS Lesion serial characteristics

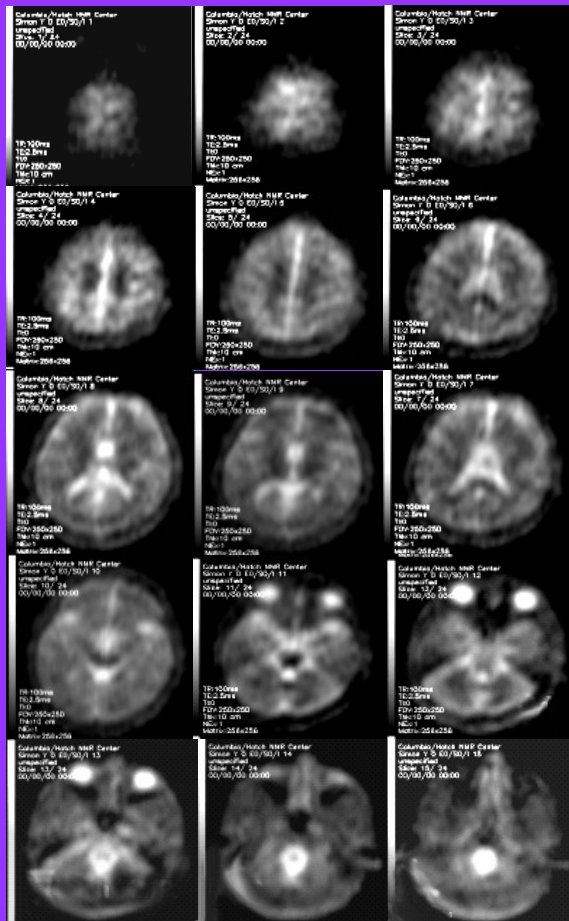
Sodium MRI Imaging

- **Brain**
- **Prostate and Breast Tumors for Chemosensitive effect**

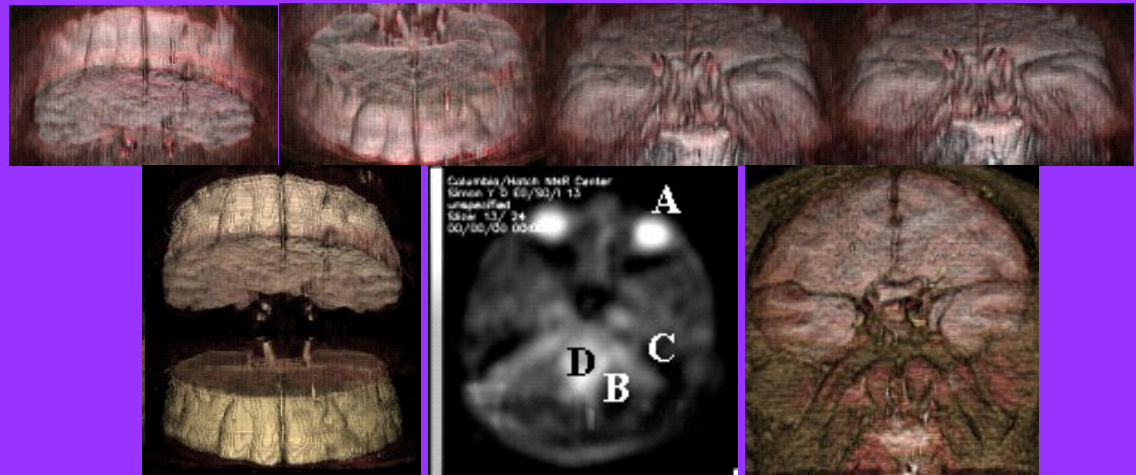
INTRODUCTION

- **SQ, DQ, TQ, MQ MRI methods measure cellular sodium**
- **Inversion Recovery(IR) pulse sequence specifically alters null point of $[Na]_i$**
- **$[Na]_i$ is elevated during apoptosis due to membrane disrupted sodium pump**
- **High 18-FDG uptake in prostate tumor is due to elevated glycolysis in tumor**

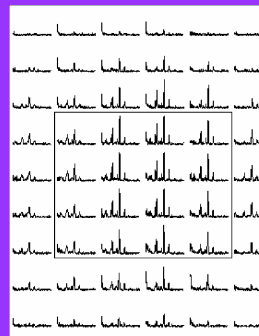
Hybrid Sodium-Proton MR spectroscopic imaging: Postprocessing



Sodium MRI images



Volume Rendered 3D Reconstructed images



Proton MRSI (left) and Sodium MRI(right)



Where we go from here?

Specific Aim 1:

Better extracellular sodium suppression using variable inversion times (TI) and phase coherence

Specific Aim 2:

Intracellular sodium concentration by alternate methods AAS, EBCT, SBF/FACT

Specific Aim 3:

Sodium gene expression and channels

Research Team

- UCSF:

- Peter Vermathen
- Andrew Maudsley
- Michael Weiner

- University of Texas
Medical School,
Houston

- P.A.Narayana
- Hong Yu
- Jerry Wolinsky
- Surjit Datta

- Columbia University &
Albert Einstein COM, NYC

- Richard Kline
- Paul Cannon
- R.K.Gupta

MRSI in Multiple sclerosis: Techniques

- **Serial studies of multiple sclerosis(MS) lesion load**
- **Serial brain metabolite imaging (Lipids)**
- **1.5 T MR chemical shift imaging(CSI)**
- **Spectral analysis**

Introduction

- **H-1 gray matter- or white matter- and CSF-suppressed MRI images visualize the MS lesion**
- **MS lesion metabolite fingerprint and volumes suggest progression of MS**
- **MRSI may be tool for drug monitoring in MS**

MRI+MRSI in MS: Technical Aspects

- **RF inhomogeneity correction**
- **3D image registration**
- **Metabolite concentrations and % tissue composition**
- **Automated segmentation of Gd-enhanced MS lesions**
- **Serial lipids and other metabolite changes in MS lesions and different regions**
- **Serial lesion volumetry**
- **GM seated MS lesions and NAGM vs NAWM**

Problem partly solved

- **Selective WM/GM suppression: (AFFIRMATIVE and QC pulse sequences)**

37,94,102(1997)

MRM

- **Lesion Volume, metabolite image-generation (APSIP and MRIAP)**

(MRM B,106,58,1995)

**AFFIRMATIVE: Attenuation of Fluid by East Inversion
Recovery with Magnetization Transfer Imaging with Variable
Echoes by FLAIR and MTC into spin echo**

QC: Quadruple contrast;

APSIP: Automated Spectroscopic Imaging Program;

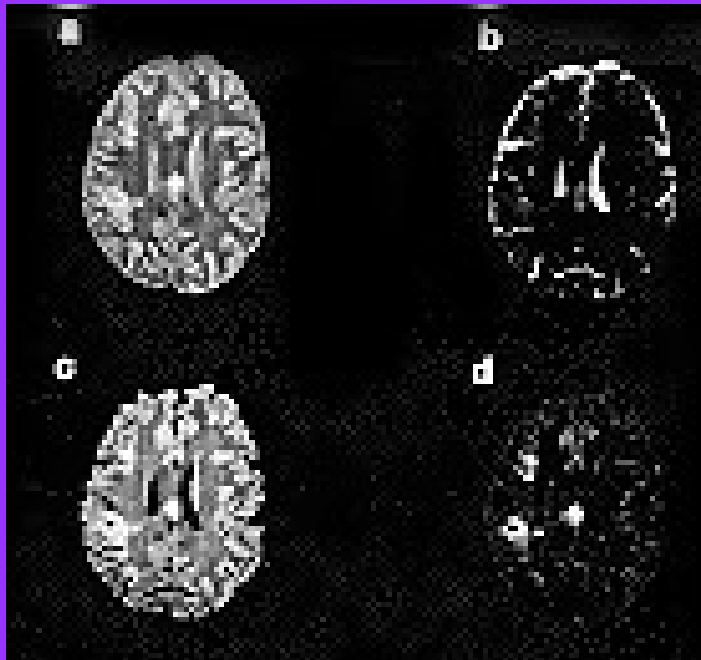
MRIAP: MR Image Analysis Program

Techniques used

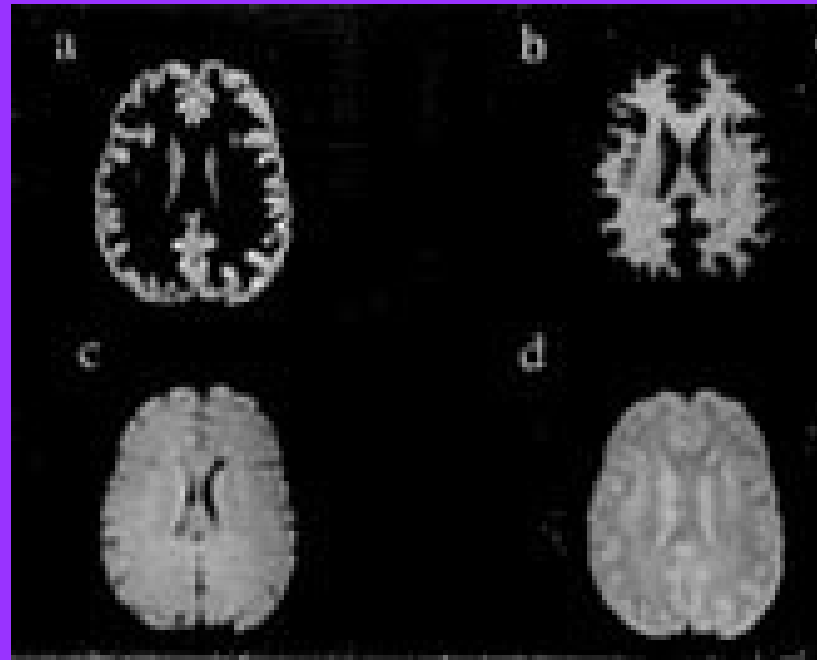
- **MR sessions for imaging: Sagittal, Axial **AFFIRMATIVE/QC**, MRSI, Pre-/post Gd enhancement**
- **Image Processing: RF correction**_{MRM 33,396(1995)} ;
- **3D-MR Image registration** _{JMRI 6,939(1996)}
- **Supervised segmentation(AFFIRMATIVE)**
- **Seed growing segmentation(QC images)**
- **Automated MS lesion segmentation** _{MRM,39,935(1998)}
- **Serial H-1 MR spectroscopic Imaging + MS lesion voluming: **VIEWER**** _{Ann Neurol,43,56(1998)}

Two approaches to enhance MS lesion contrast

- AFFIRMATIVE

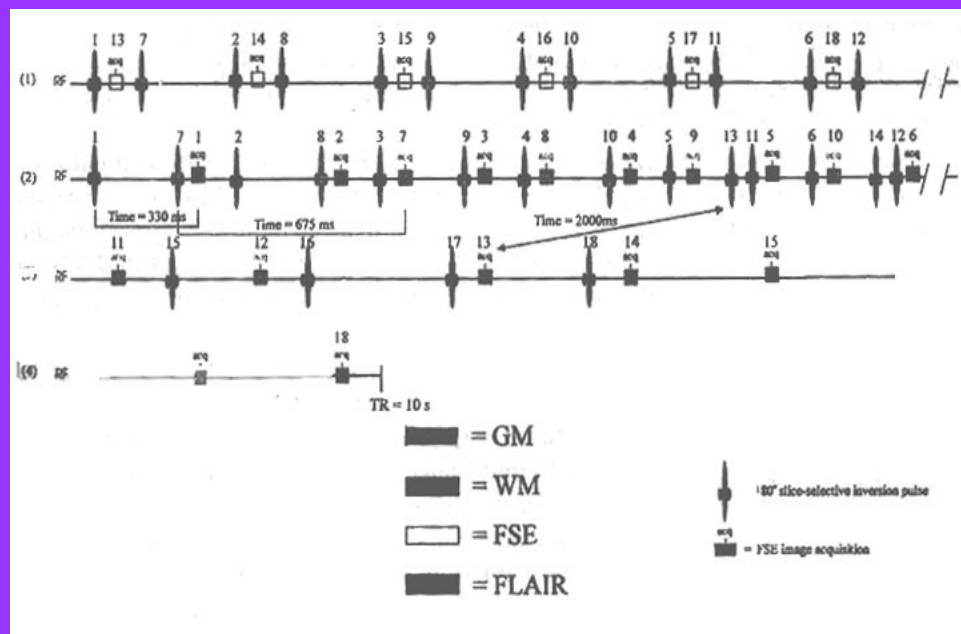
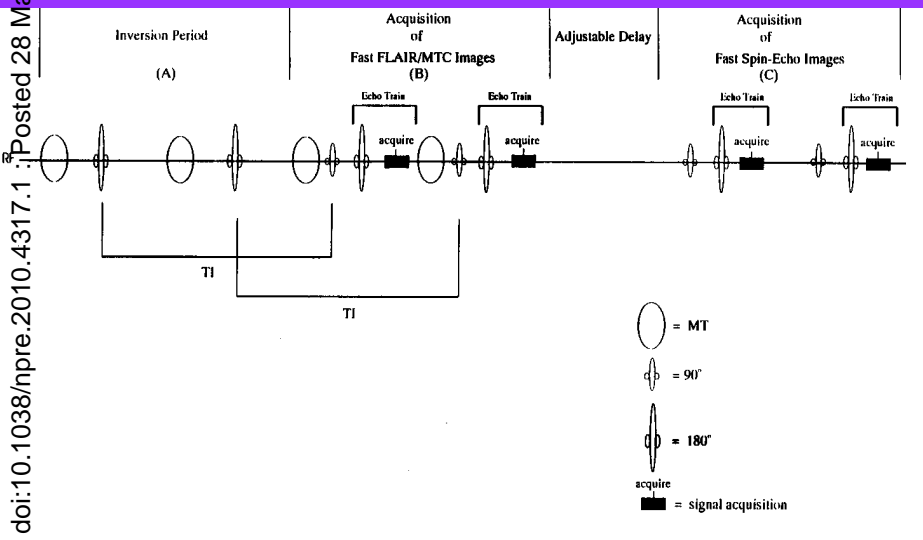


Quadruple Contrast



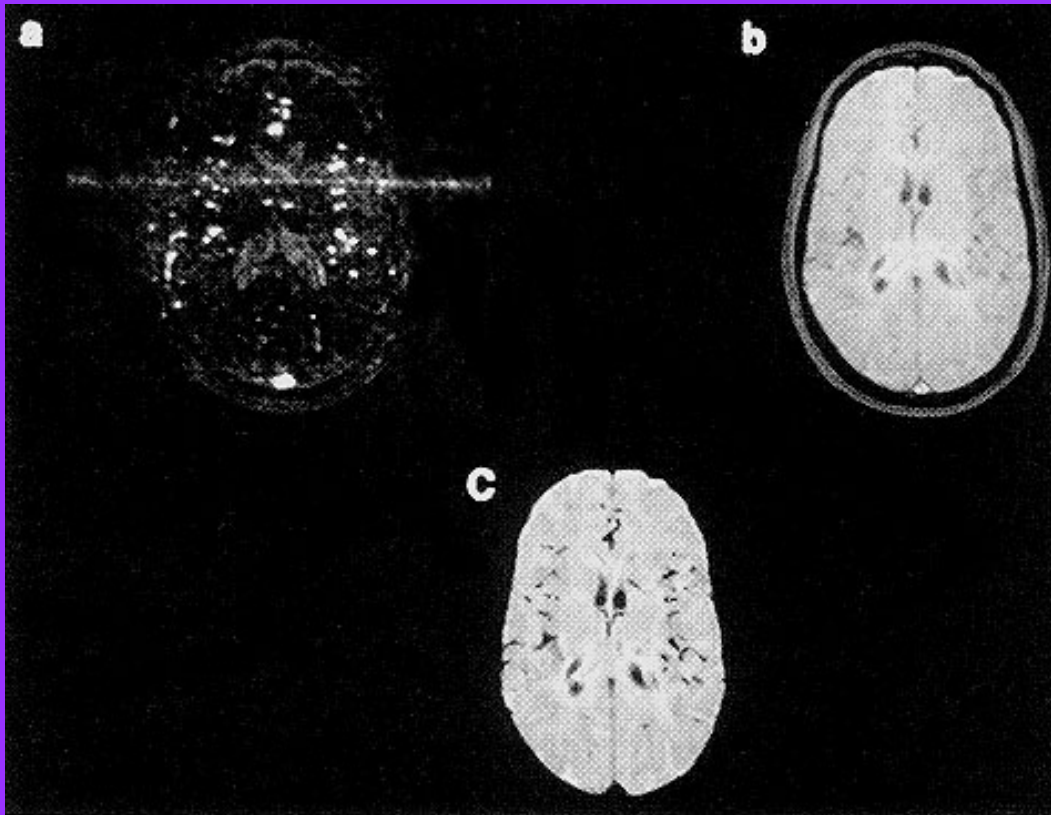
AF FIR MATI VE Pulse Sequence and Quadruple Pulse Sequence

Nature Precedings : doi:10.1038/npre.2010.4317.1 v1 Posted 28 Mar 2010



- Left: Fast SE + MT pulse at early and late echoes
- Right: 2 inversion pulses for selective suppression of GM, WM, CSF fractions

Phase-contrast Flow Images

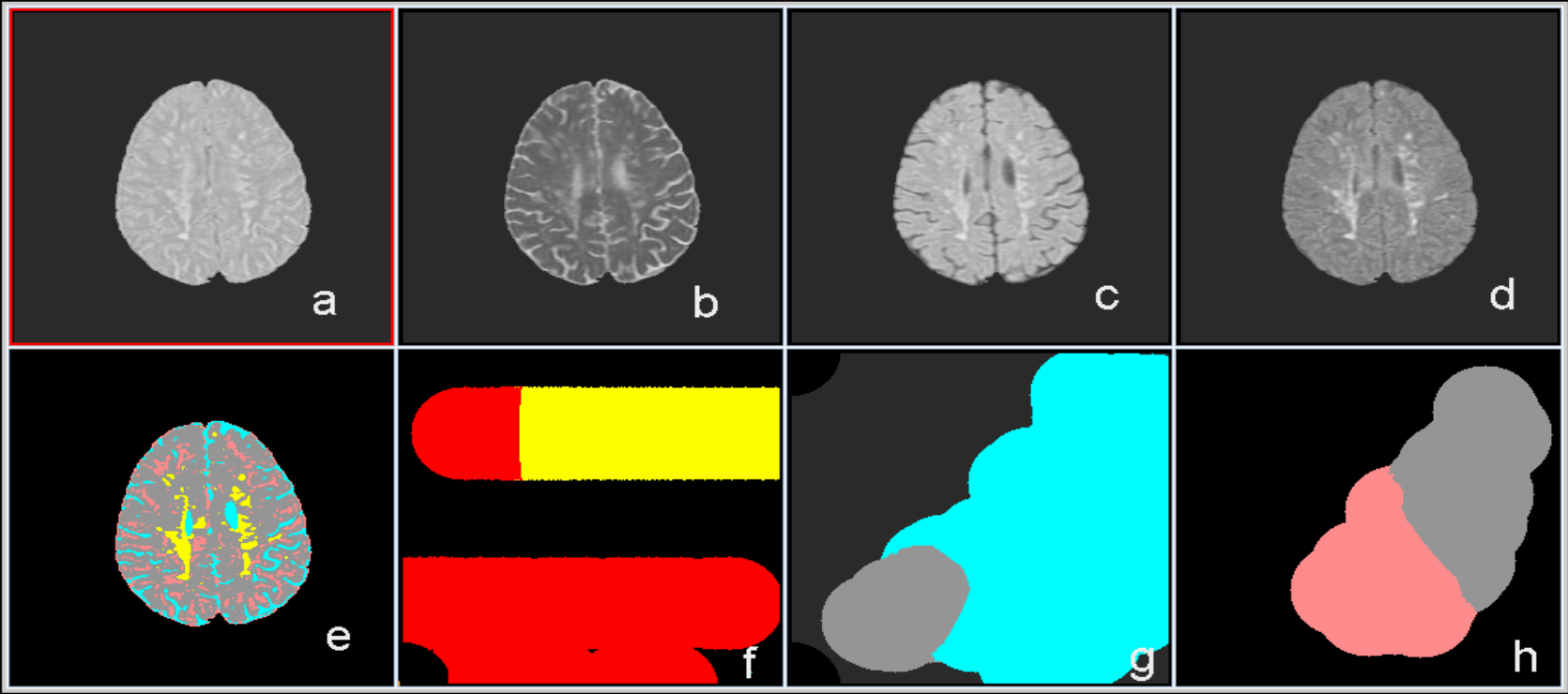


- MRA image (a)
- Magnitude Image(b)
- Flow Image (c) without scalp
- Cerebro-vascular spins are visible (a)

Magnetic Resonance Image Analysis Package(MRIAP)

- **Image Conversion**
- **Read/Modify Header**
- **Image Processing:**
 - **Anisotropic Diffusion Filter**
 - **Manual Editing:**
 - **Connectivity,line drawing, island removal,intensity thresholding, striping remained pixels**
 - **Automated Slice Editor**
 - **RF inhomogeneity correction**
 - **Flow Generation Image Generation**
 - **Image Set Recording**
 - **MR Angiography**
- **3D Brain Registration**
- **Image Segmentation:**
 - Seed growth/ manual editing**
 - Supervised Segmentation: 2D feature**
 - map/multiple feature map segmentaion
 - Automatic Segmentation**
 - Lesion Contrast Enhancement**
- **Volumetric Analysis:**
 - 2D interactive**
 - 3D Automated lesion volumetry**
- **Multiple Display: SID**

Magnetic Resonance Image Analysis Package



2D Feature Map Generation and Setup

| | File Path | Display Window | Feature Window Axis #1 | Feature Window Axis #2 | Intensity Range Axis #1 | Intensity Range Axis #2 | | |
|-----------------|-----------------------------------|----------------|------------------------|------------------------|-------------------------|-------------------------|------|------|
| Feature Map #1: | /opt/mr/lib/segmaps/lesion320.map | 6 7 8 | 3 | 4 | 1024 | 1024 | Load | Save |
| Feature Map #2: | /opt/mr/lib/segmaps/csf.map | 6 7 8 | 2 | 3 | 1024 | 1024 | Load | Save |
| Feature Map #3: | /opt/mr/lib/segmaps/tissue.map | 6 7 8 | 1 | 3 | 1024 | 512 | Load | Save |

Image Segmentation Canvas Control

Window #1: Display Echo #: 1 2 3 4

Window #2: Display Echo #: 1 2 3 4

Window #3: Display Echo #: 1 2 3 4

Window #4: Display Echo #: 1 2 3 4

Window #5: Display Echo #: 1 2 3 4

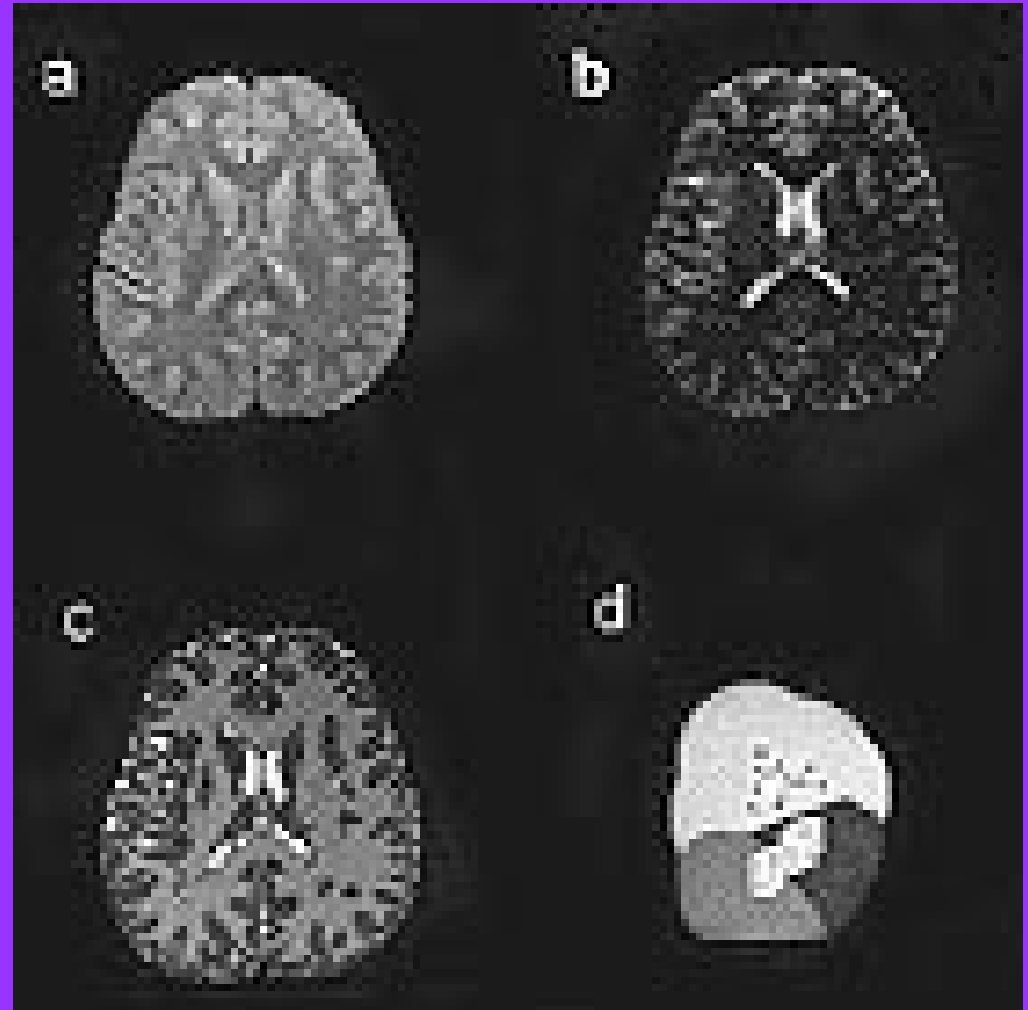
Image Segmentation Main Control

Segmentation Mode:

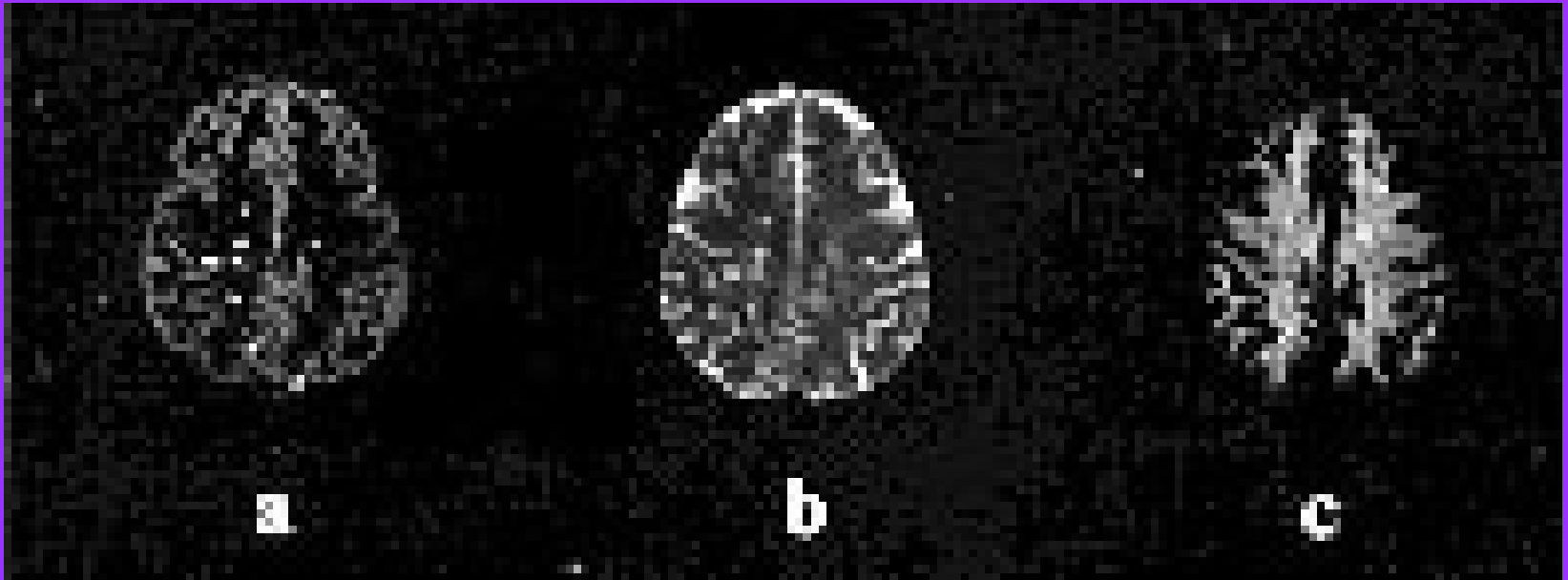
- Manual -- Paintbrush
- Manual -- Seed Growing
- Manual -- Editing
- Manual -- Erasing
- Supervised
- Automatic
- Contrast Enhancements

RF inhomogeneity Correction: Tissue Cluster Analysis

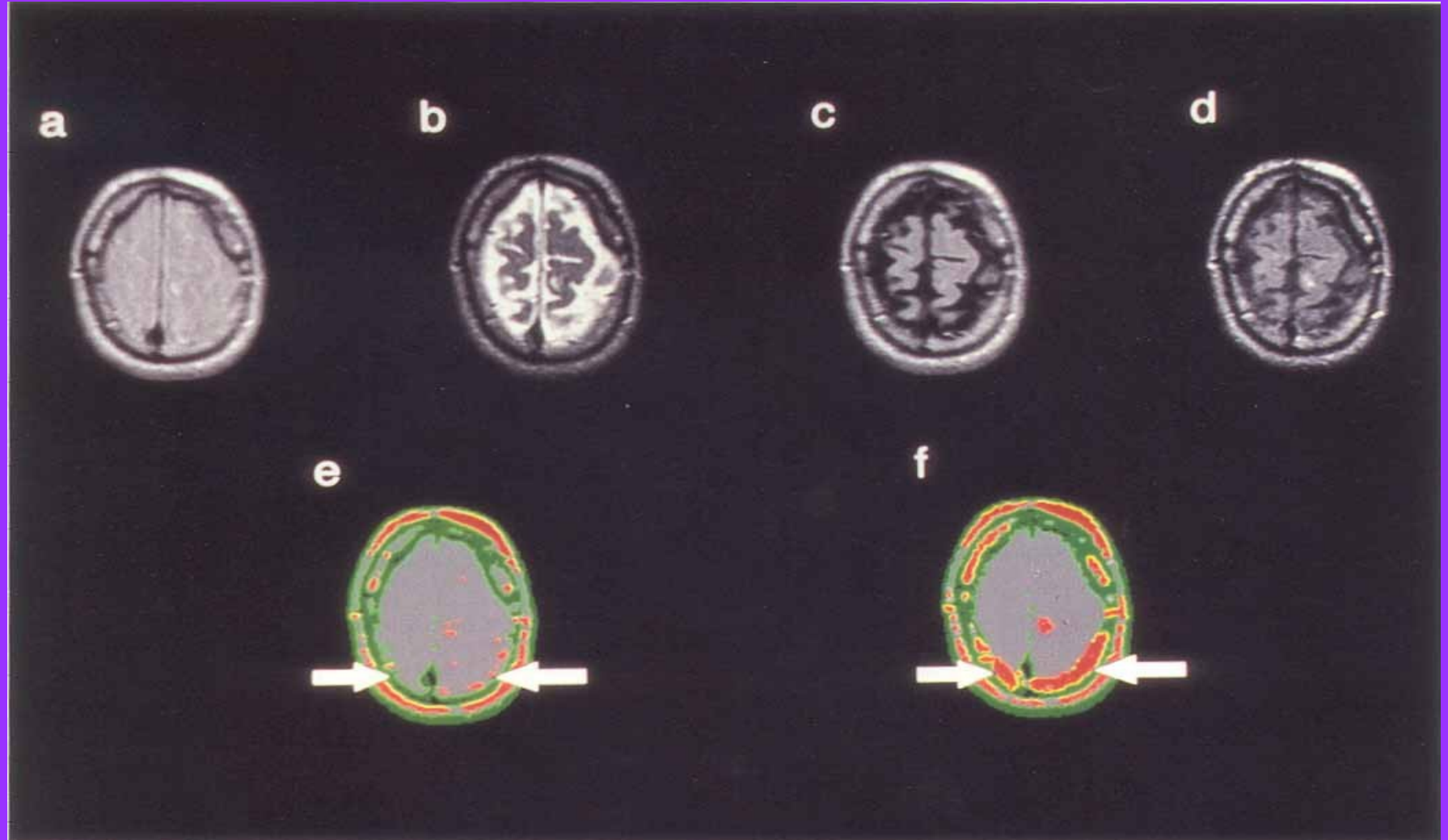
Normal axial brain images at TE=16 ms(a);
TE=80 ms(b);
Segmented image(c);
Tight clustering of feature map (d)



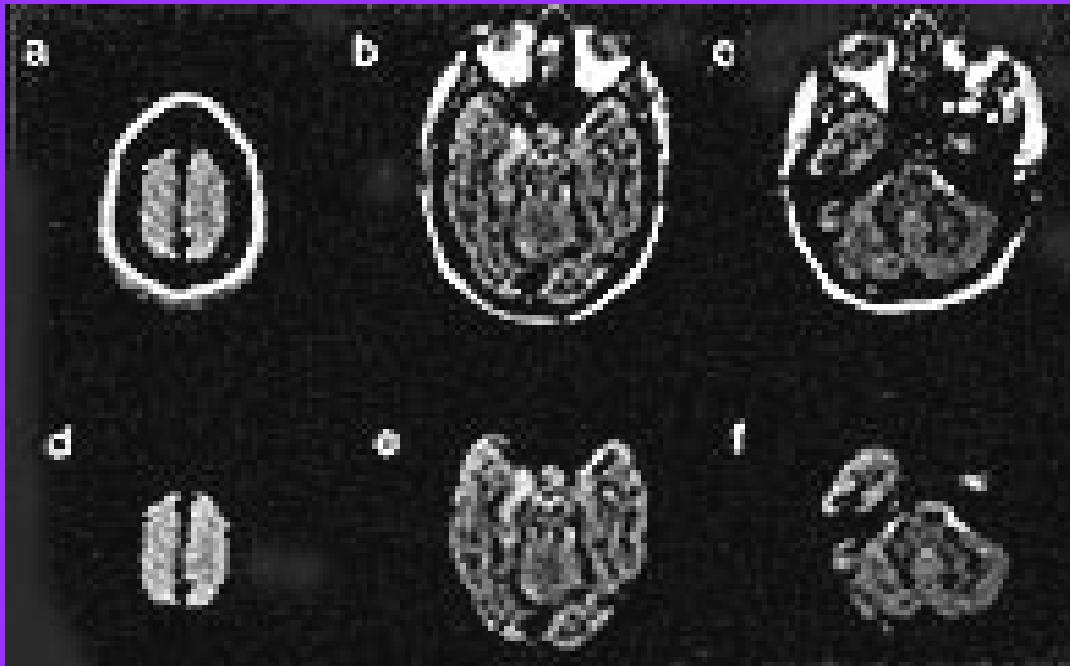
Selective Tissue Suppression Method



Extrameningial Tissue Removal by Connectivity Algorithm



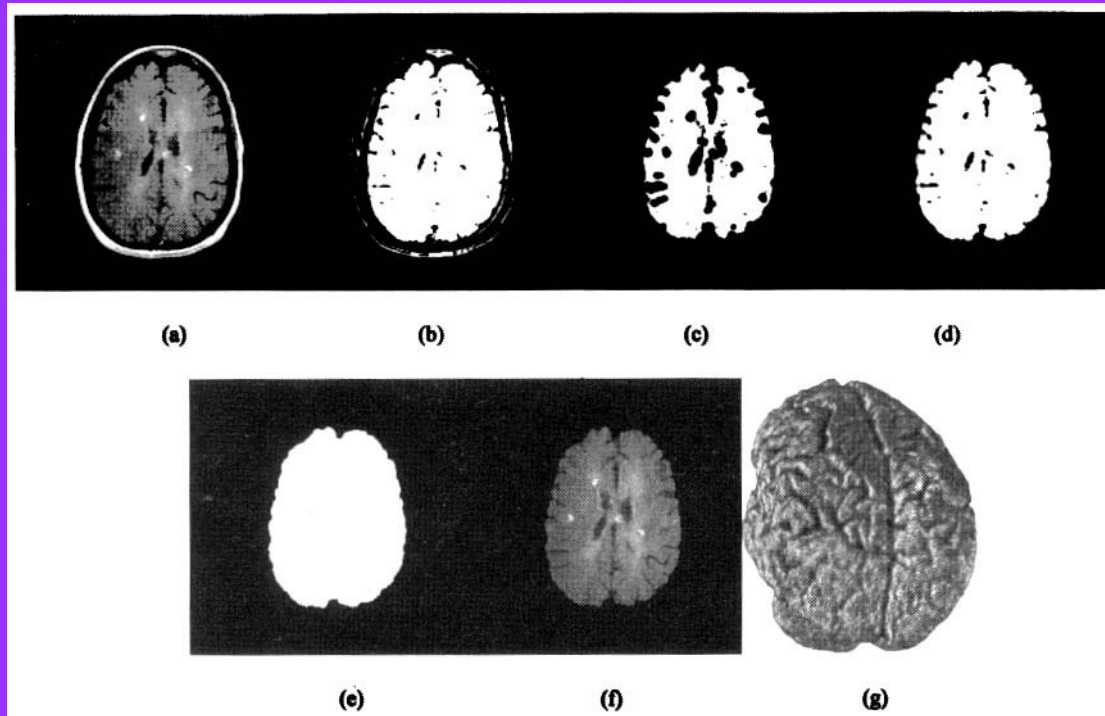
MR Images Prior to And Following Removal of Extrameningial Tissue



- **Multispectral, nonparametric Perzen window classifies Extrameningial Tissue**

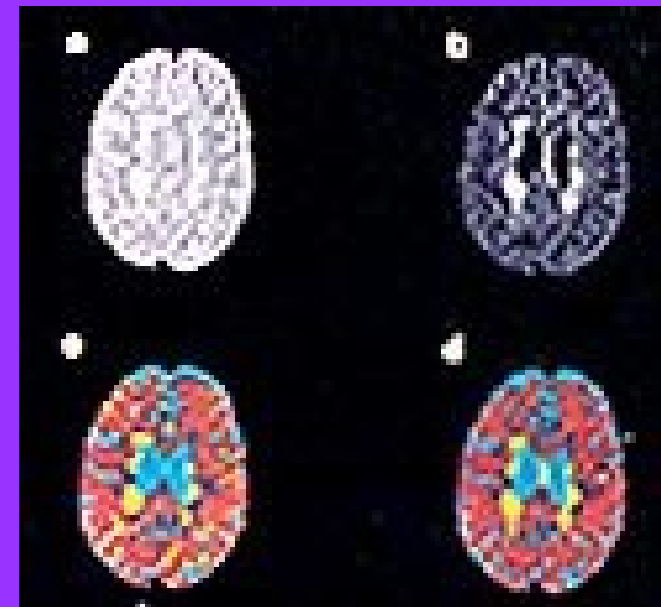
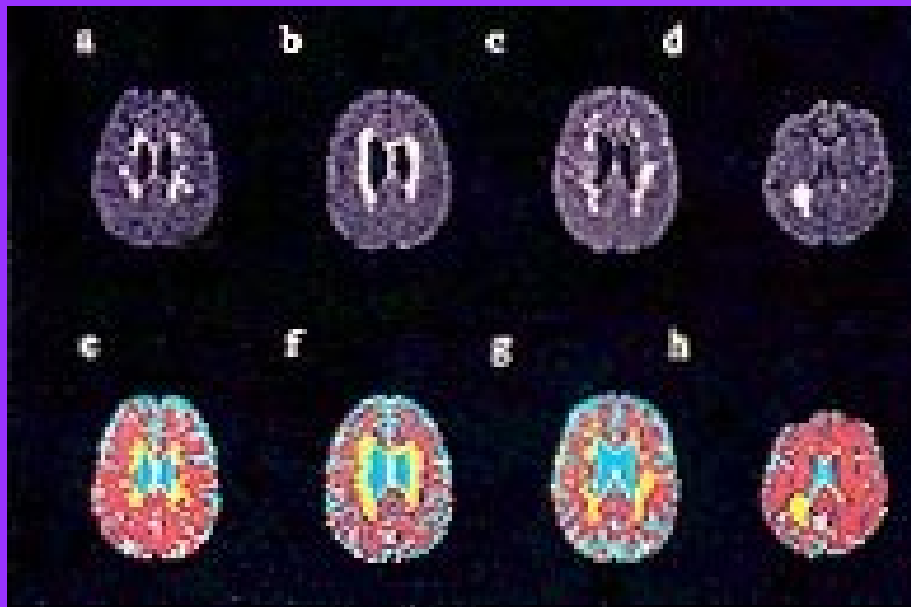
- **Fluid attenuation and magnetization transfer contrast**

Segmentation in Brain



**a.Raw image;b.Thresholding; c.Erosion;d.Dilatation; e.Closing
f and g.Volume rendered images**

Automated Tissue Segmentation (Perzen Non-parametric Method)

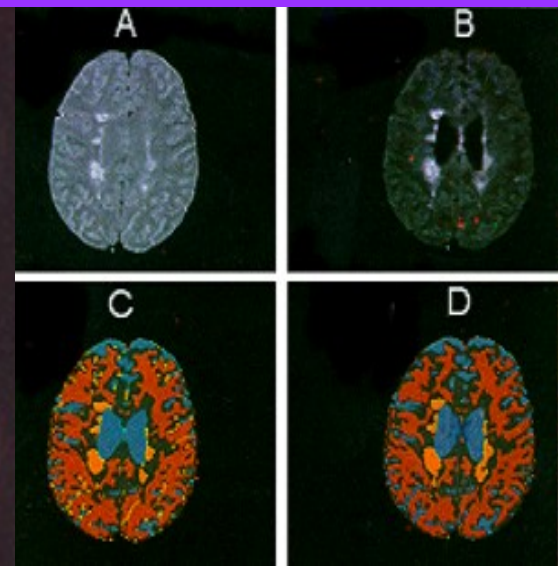
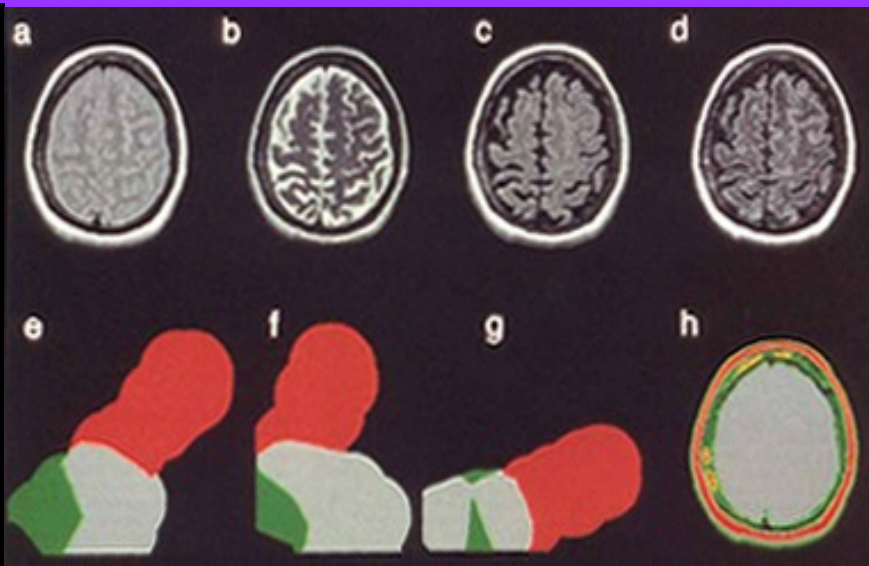
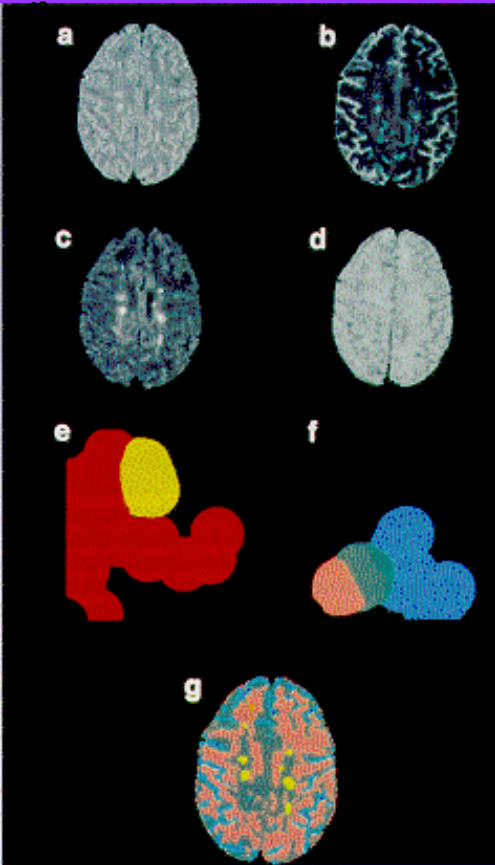


- **MRI images at different level (top)**
- **Different color-coded tissues(GM,WM,CSF, lesion and non-lesion) by supervised training data-set**

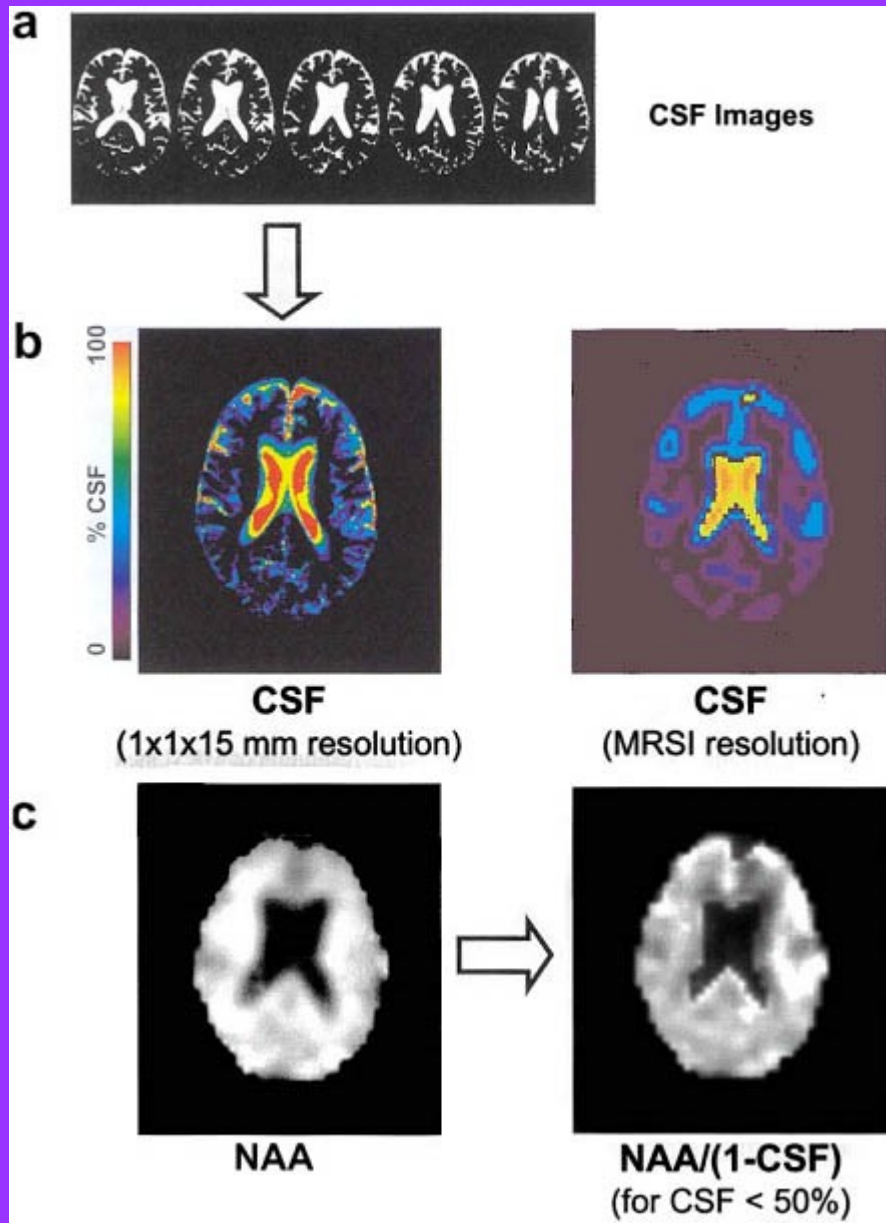
Non-Parametric Segmentation: (Perzen

Window Feature Map Generation)

Mar 2010



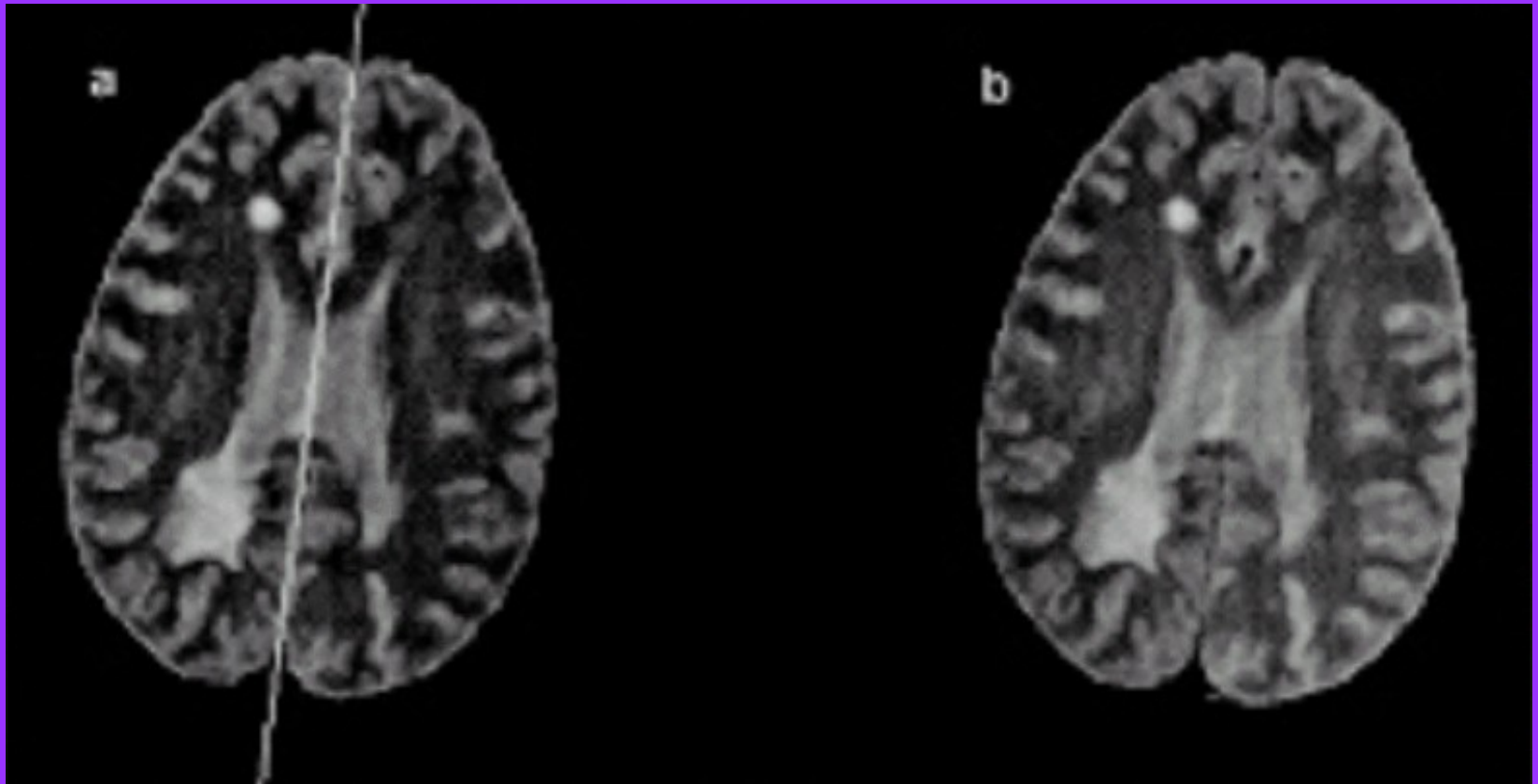
- AFFIRMATIVE Images and feature maps(Left)
- Quadruple contrast images and feature maps(on right)
- Dual Approach of false lesion minimization



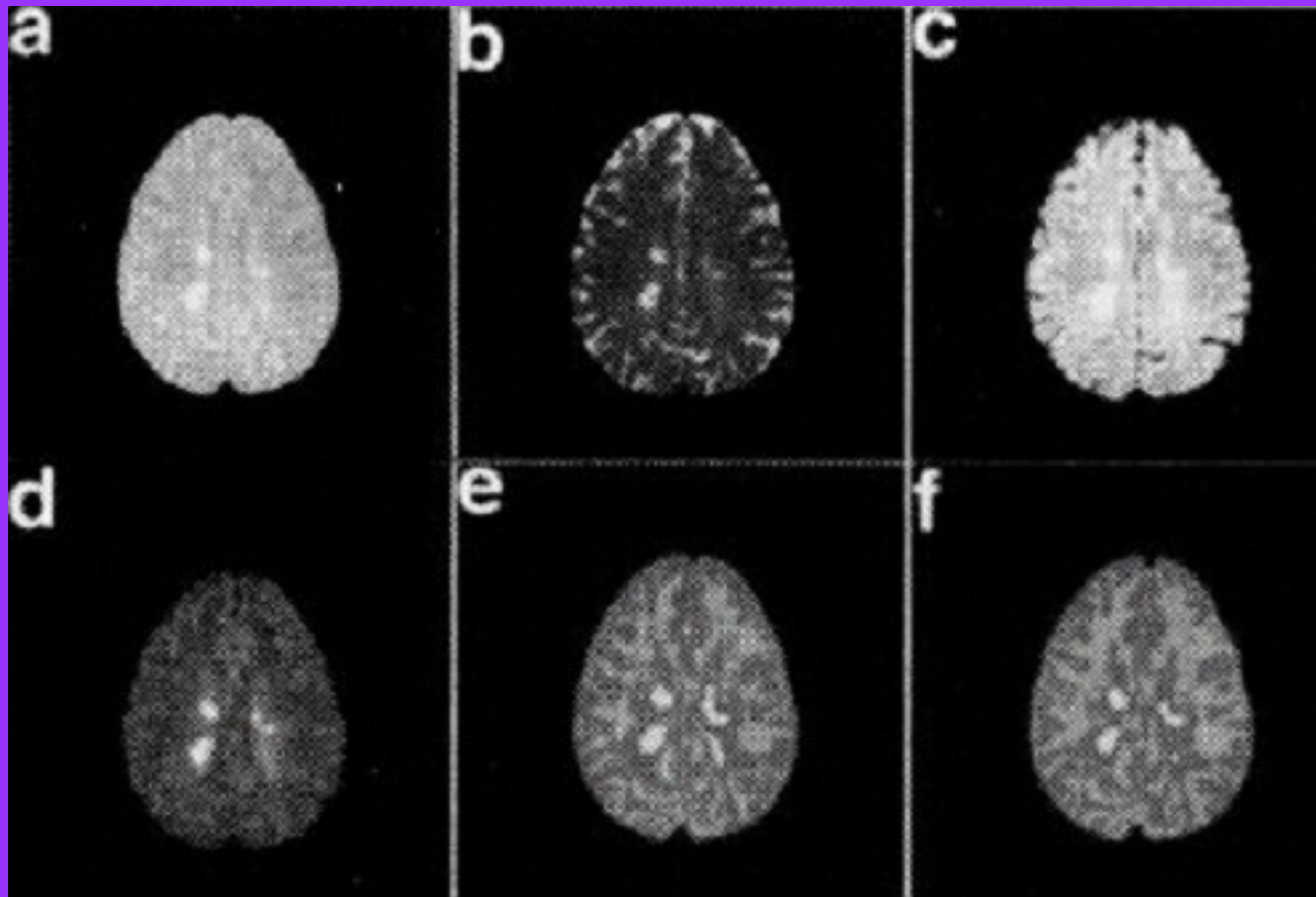
Effect of CSF Nulling

- CSF Minimization
- Atrophy Correction for NAA

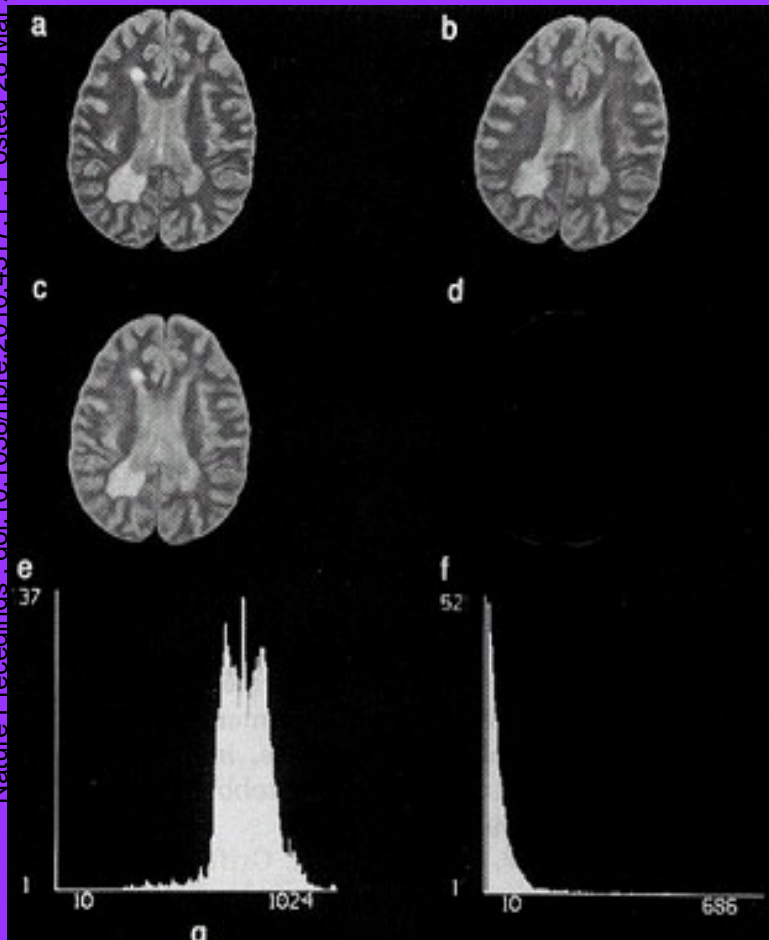
Registration: Interhemispheric Fissure Autosearch Method



The Effect of Histogram Normalization on Segmentation

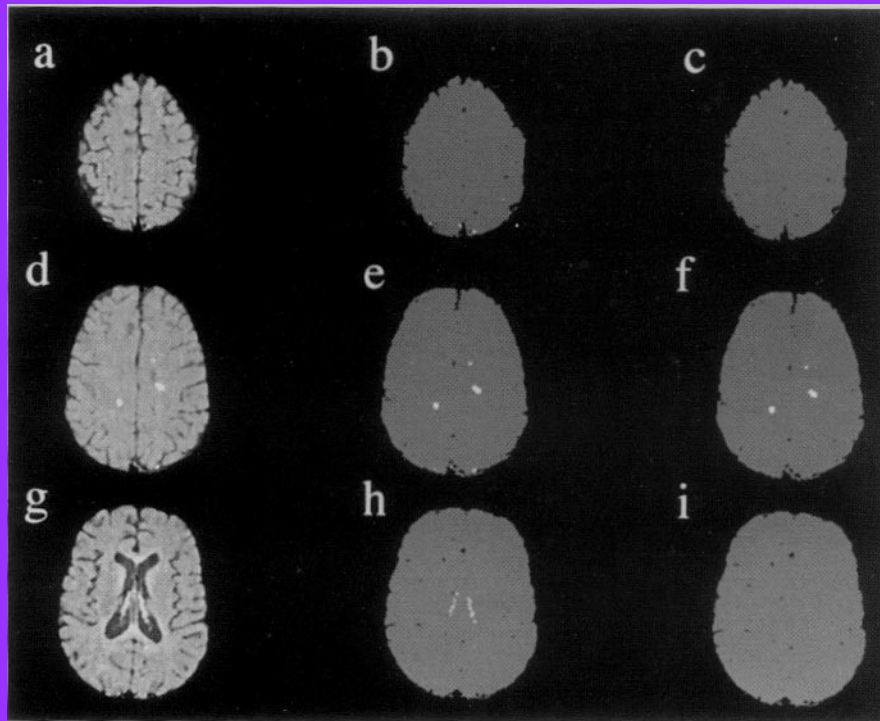


3D Registration and histogram analysis



- a. Axial T1 image of MS(a)
- b. Computer Generated offsets at $5^\circ \times 3^\circ \times 2^\circ$ rotation in z,y,x axes(b)
- c. Registered image(c)
- d. Subtracted image(d)
- e. Histogram pixel value in image(e)
- f. Histogram pixel value in image (f)

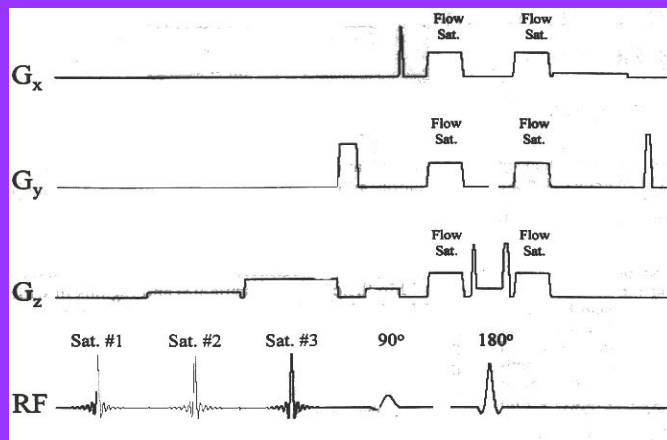
MS Lesion Gd-Enhancement



- Pre-contrast Images (1st column)

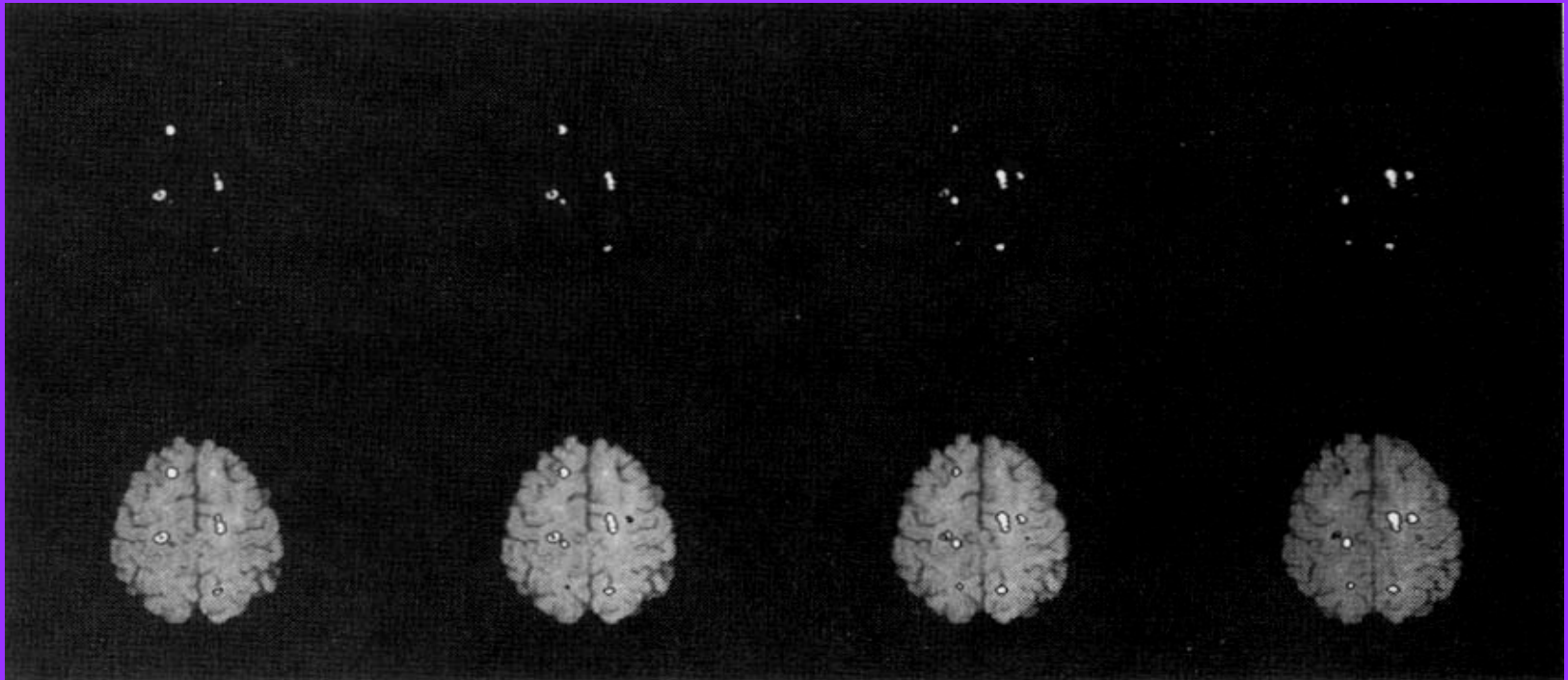
- Post-contrast (false positive 2nd column)

- Post-contrast (corrected 3rd column)



- Method: Saturation pulses and gradient dephasing Trapezoidal gradient waveforms select Gd-enhanced pixels.

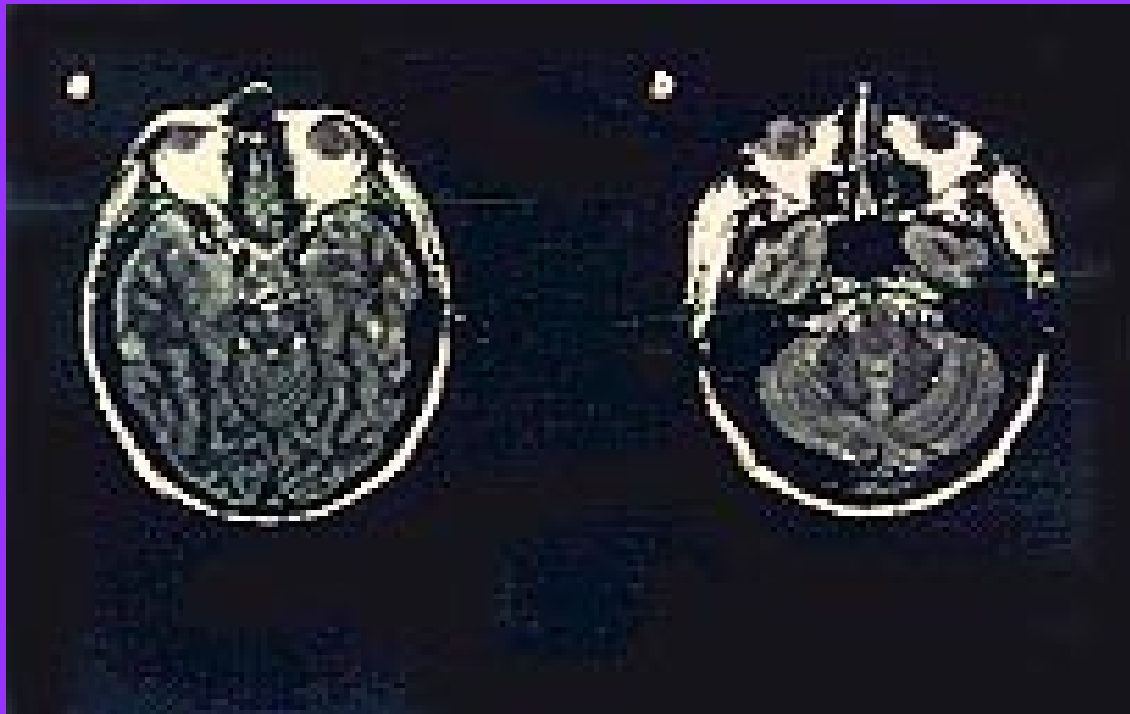
Delineation of Gd-contrast enhanced MS lesions



Top Row: GM+WM+CSF suppressed images

Bottom Row: Delineation of MS lesions

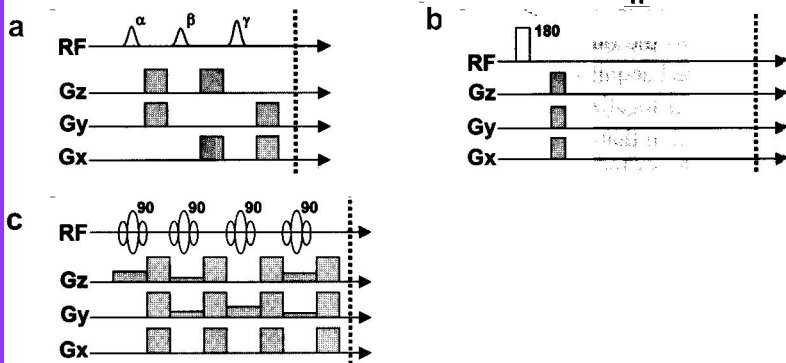
Connectivity Algorithm and Removal of Extrameningial Tissue



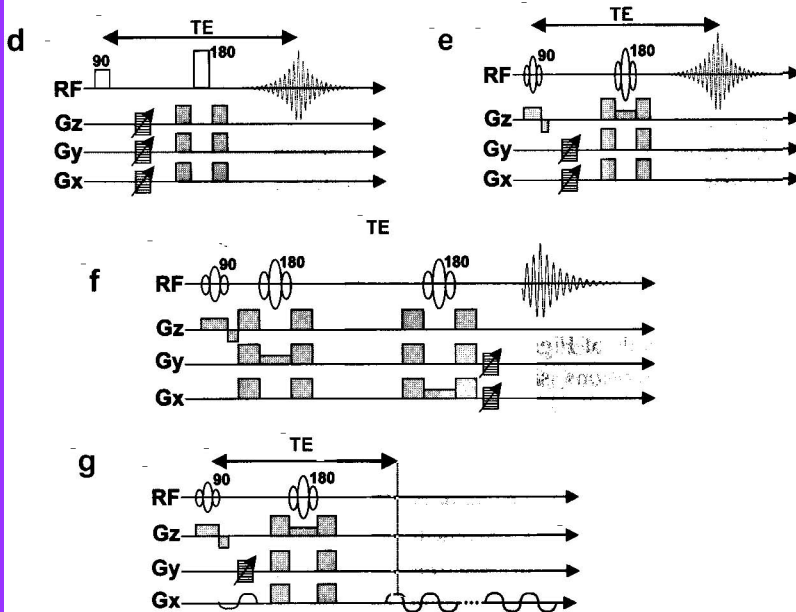
- Green lines represent the selective pixels separating out brain regions from optical nerve and other parts

MRSI Pulse Sequences

Preparatory Sequences

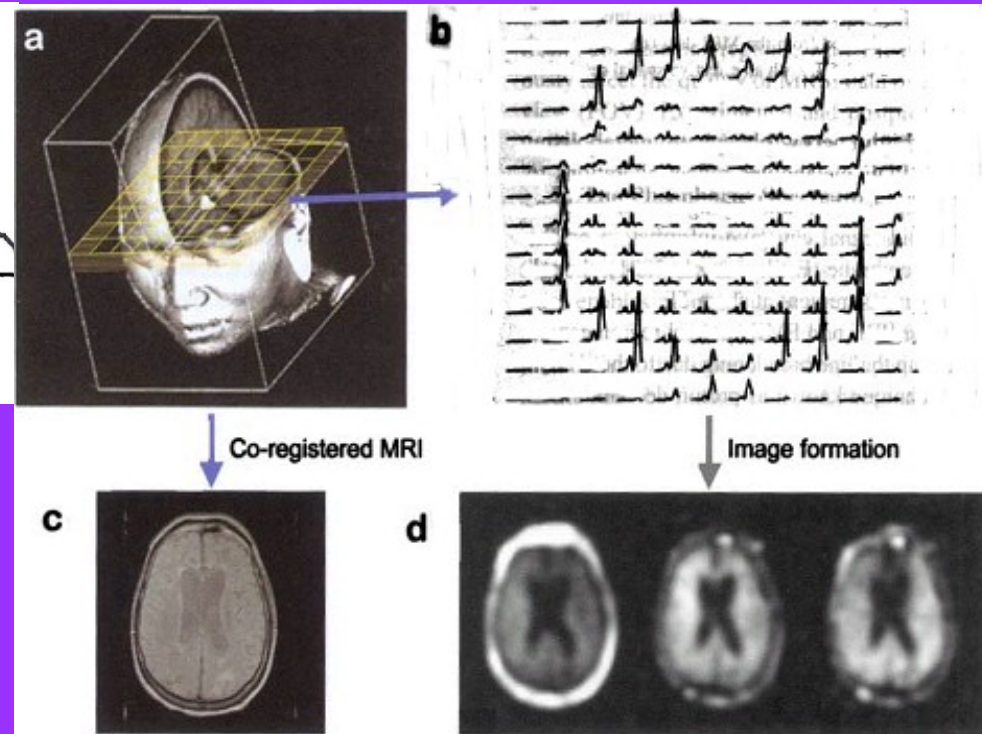
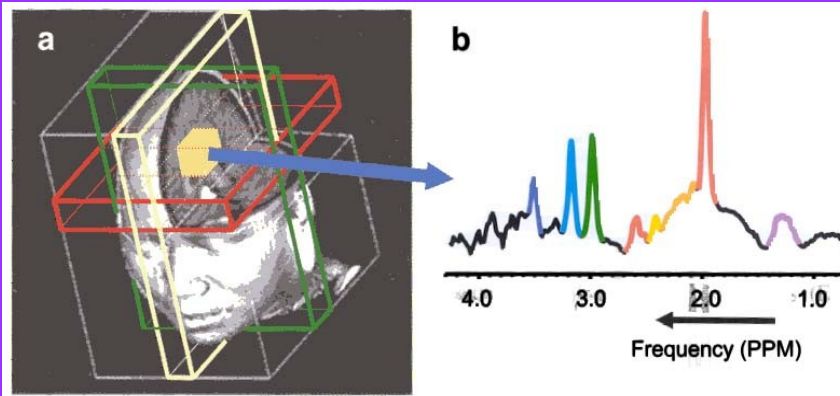


MRSI Sequences



- 90° and 180° RF Pulses applied with variable gradient sets
- Spectroscopic voxels are selected by use of fat saturation bands
- Protons of -CH₃, -CH₂ etc show chemical shift

MRSI Slice Selection: Metabolite Maps and CSI imaging



- Selective Single Voxel Spectroscopy (left)
- Single Plane Chemical Shift Imaging

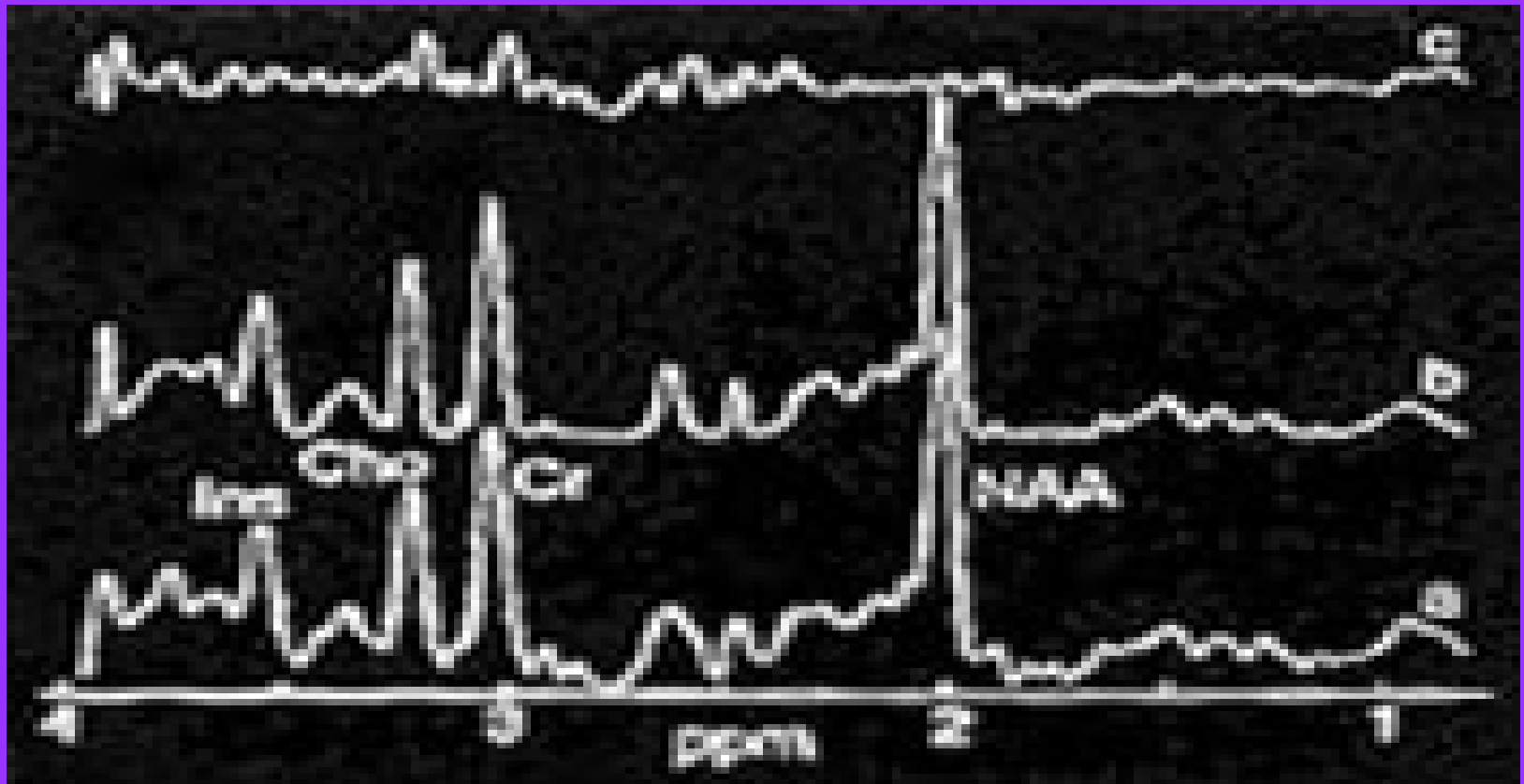
Techniques Used in MRSI (APSIP package)

- For H-1 Spectral model library for spectral simulation a priori information;
- Denoising algorithm for baseline smoothening
- Automated MRSI spectral fitting method:
 - Zero-filling, Object masking; Frequency shift(cross-correlation); iterative baseline and metabolite optimization(VOI), curve fitting

Spectroscopic Imaging and peaks: Data Processing

- **Spectroscopic VOI synthesis:**
 - masking “Octagonal OVS mask”
 - scanner co-ordinates selected
 - Pixel thickness and threshold fixed
 - Gray scale/color
 - Grid and shift
- **Data Processing:**
 - APSIP(automated processing by SI program)
 - Steps: zero filling, DC baseline, spatial apodization, CSI 1D rotate, CSI flip, CSI 2D rotate, Bo autophase, water suppression, deconvolution coefficient, time apodization, FFT

Automated Processing for Spectroscopic Imaging Package



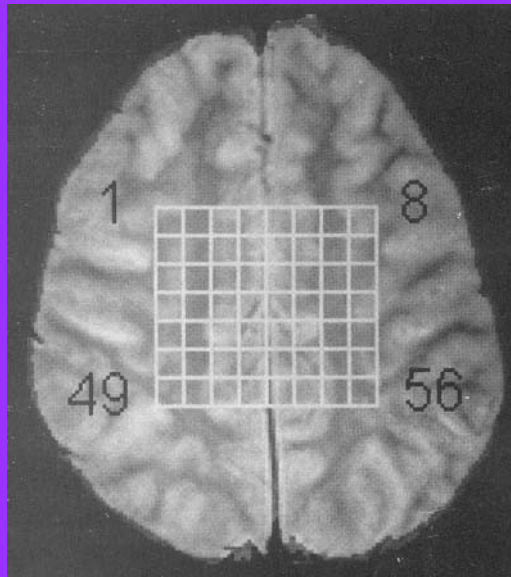
Autopsy MS lesion NMR spectral peaks

| Chemical Shift (ppm) | Assignment Metabolite | Normal WM | MS lesion |
|----------------------|-------------------------------------|-----------|-----------|
| 0.8 | lipid(-CH ₂) | | |
| 0.25 | | | |
| 1.00 | lipid(-CH ₃) | | 0.68 |
| 1.20 | triglycerides | | 1.88 |
| 1.33 | lactate | | 2.25 |
| 1.4 | alanine | | 1.80 |
| 1.5-1.55 | leucine | | 2.5 |
| 1.6 | ?? | | |
| 1.7 | glutathione ?? | | 0.6 |
| 1.8 | GABA | | 0.5 |
| 1.9 | acetate | | -- |
| 2.02 | N-acetyl aspartate (NAA) | 10.8 | 6.3 |
| 2.1 | NAA Glutamine (β methylene) | | - |
| 2.2,2.3,2.4 | Glutamine/glutamate | | 5.5 |
| 2.7 | Aspartate (β methylene) | | 1.3 |
| 3.0 | creatine/phosphocreatine(=N.methyl) | 8.0 | 6.8 |
| 3.2 | choline(-N(Methyl)3) | 3.2 | 7.2 |
| 3.4 | taurine(=N.methylene) | 0.4 | 0.45 |
| 3.6 | myo-inositol | 3.2 | 3.5 |

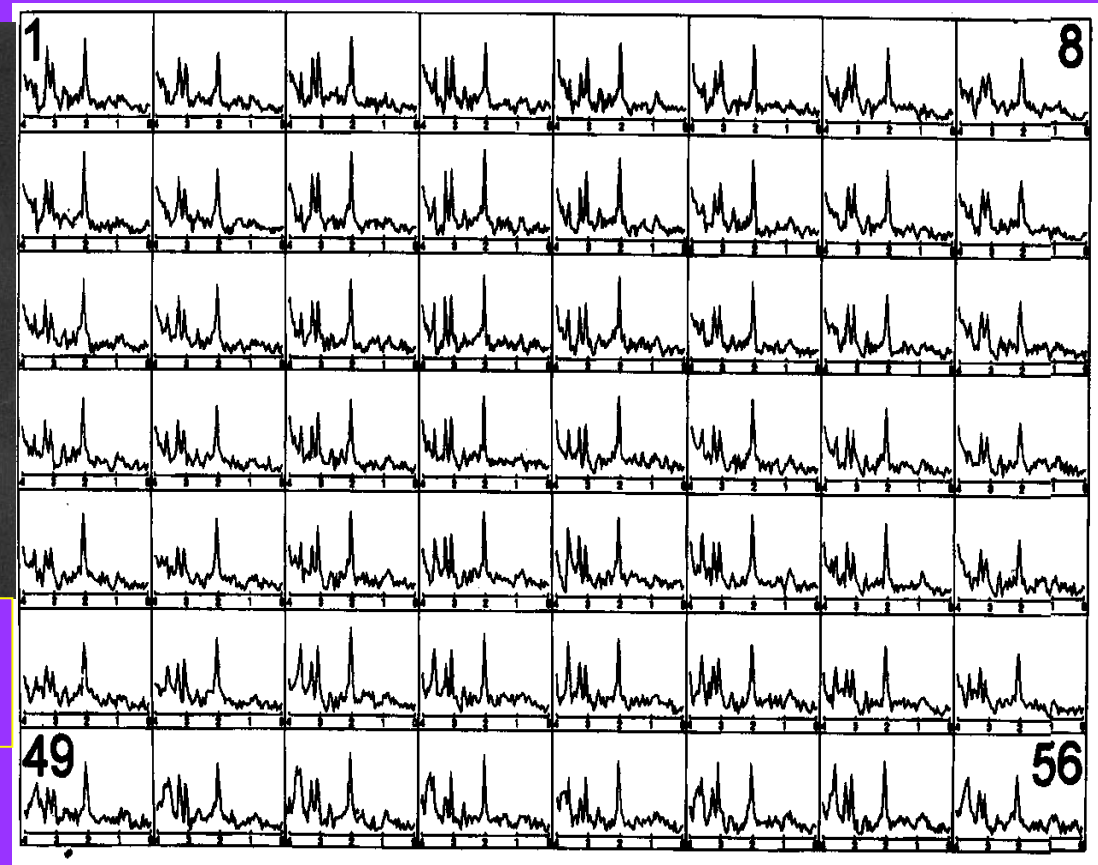
Objectives

- **Serial longitudinal studies of MS:**
 - lesion volumes**
 - Metabolites(NAA, Cr, Choline and lipids) at lesion, GM, WM sites**
 - Tissue(GM, WM) fractions**
 - Relationship with neuropsychological tests (MMSE and EDSS)**

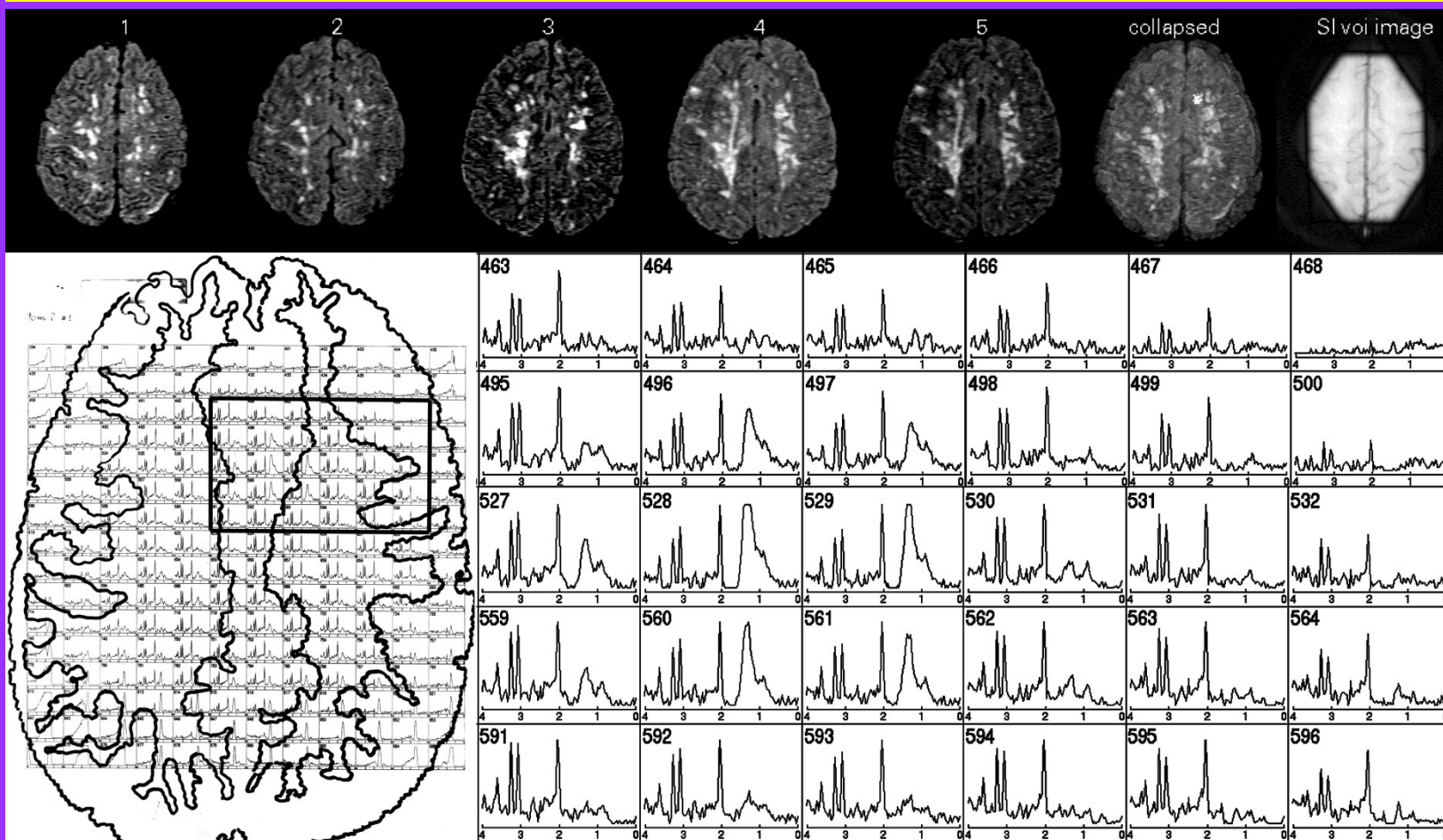
MRS metabolite peaks in Normal Volunteer



•NAA, Cr and Cho peaks visible

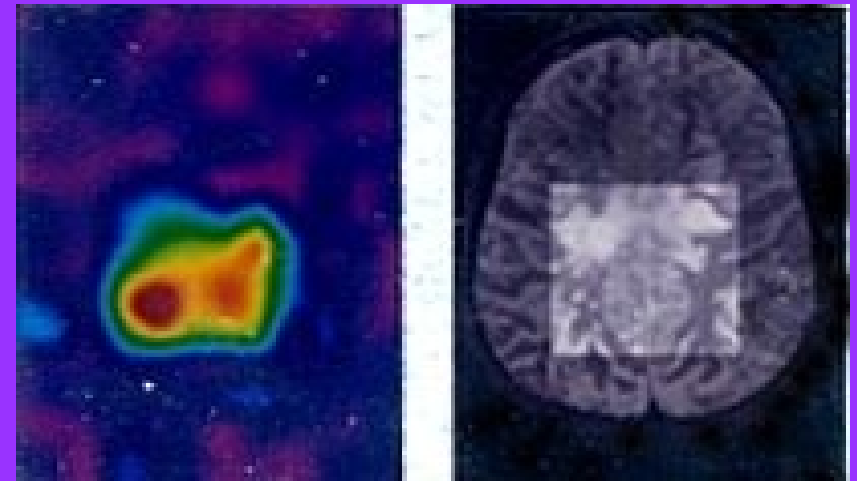
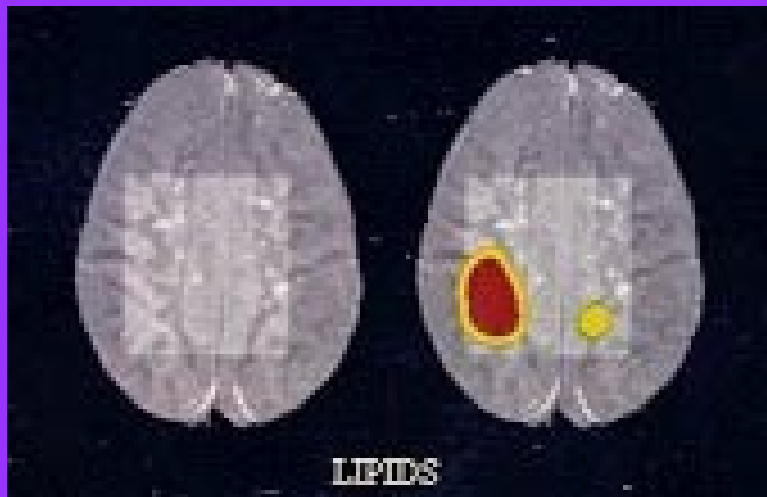


MS Lesions Showing Abnormal Metabolites in GM and NAGM After Post-Gd Contrast

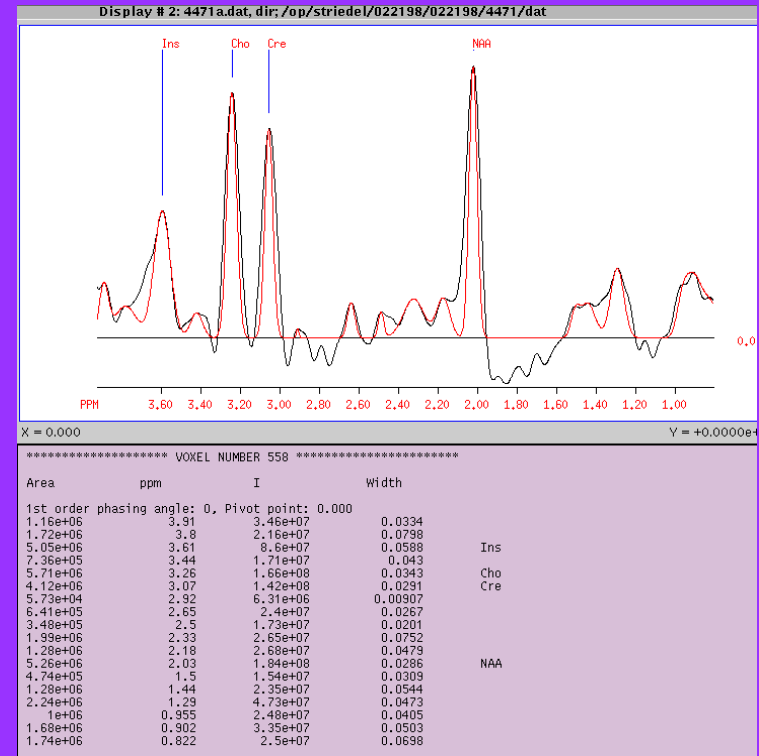
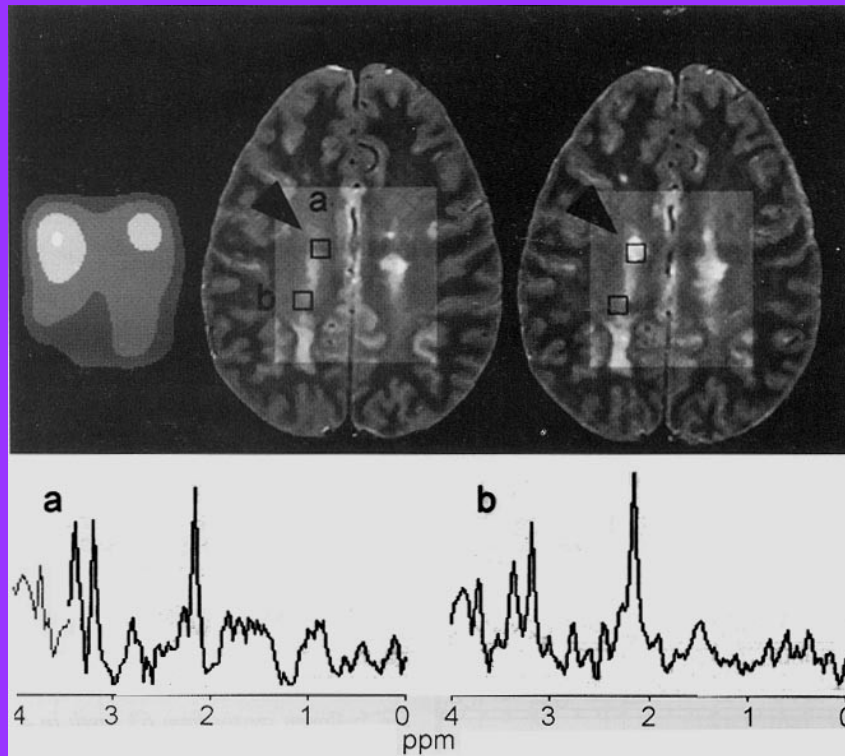


Lipids in MRI-defined MS lesions

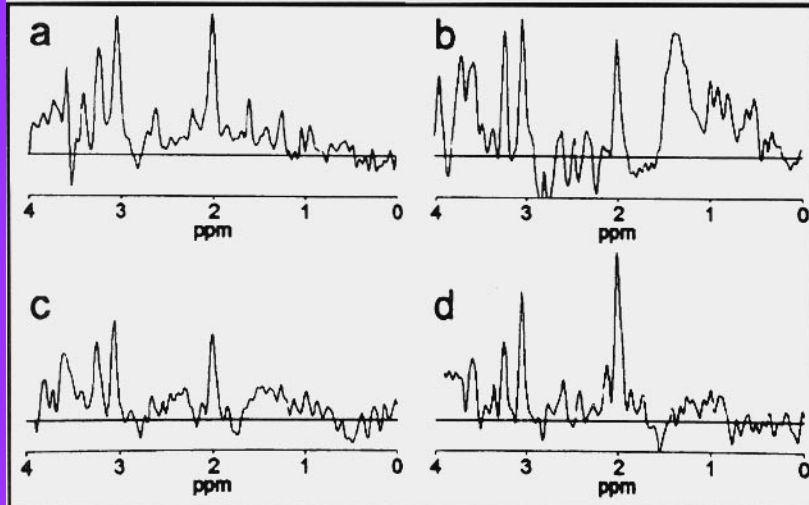
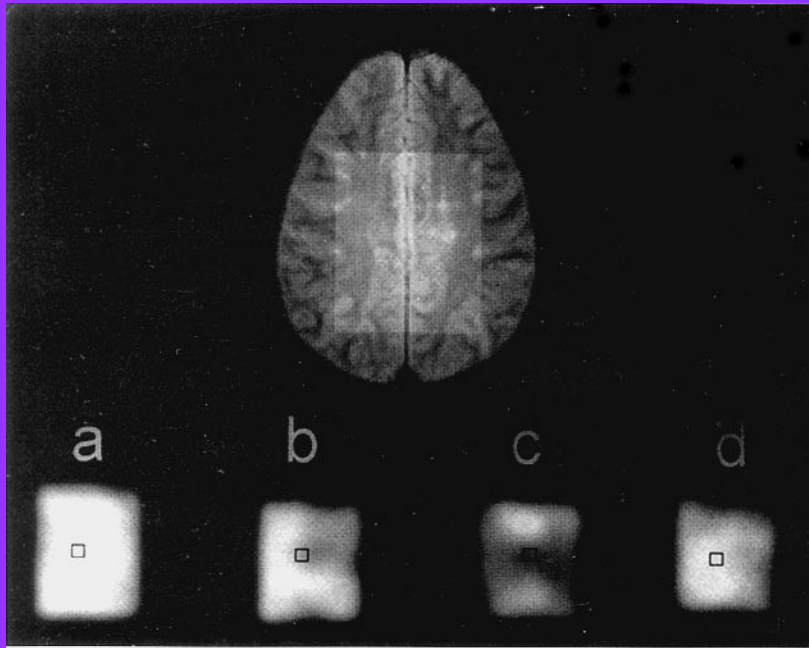
- MRI defined MS lesions
- Lipid metabolic maps (color-coded)
- Lateralization by lipids



Serial MS study: choline peaks



- Choline as inflammatory marker(perivenous inflammation)
- Related other lipids, amino-acids peaks



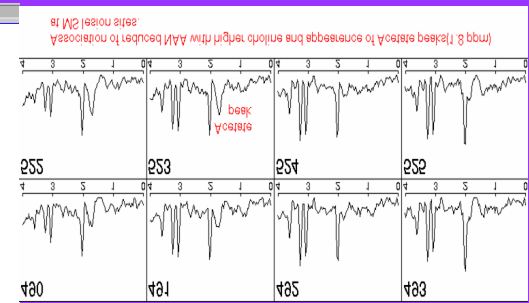
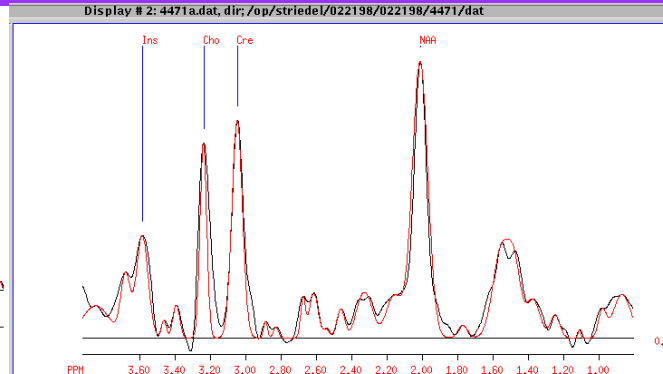
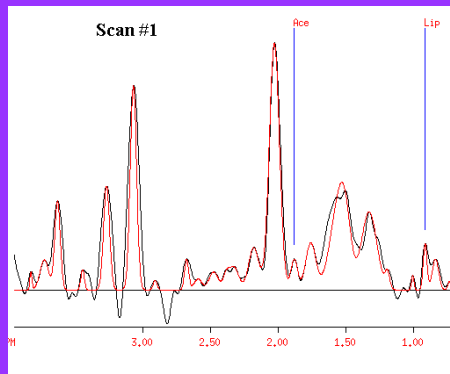
Serial Lipids in NAWM

- No MRI-defined MS lesion
- Lipids as singlet and doublet peaks

Where we go from here?

- Better spectral MRSI resolution
- New MR visible metabolite GABA, glutathione
- Amyloid proteins and CSF proteins
- Dementia classification and metabolite regional differences
- Source of NAA origin in different brain regions
- Gene expression and regional metabolite
NAA/Cr+Cho ratio
- Relationship of neurochemicals with biochemical markers (Acetylcholine esterase)

Difficulty in Spectroscopy Peak Interpretation



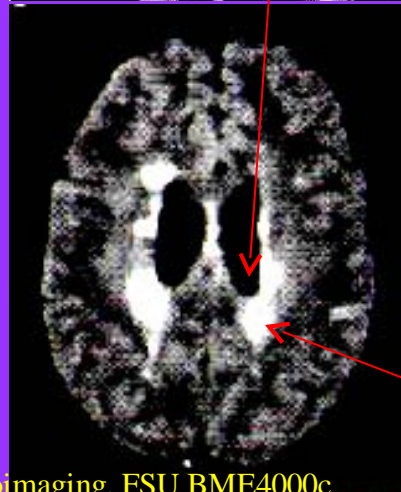
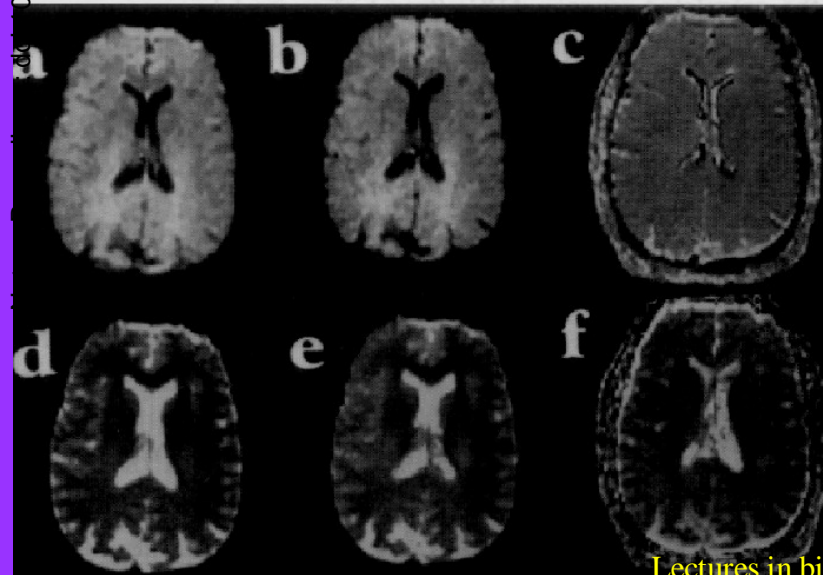
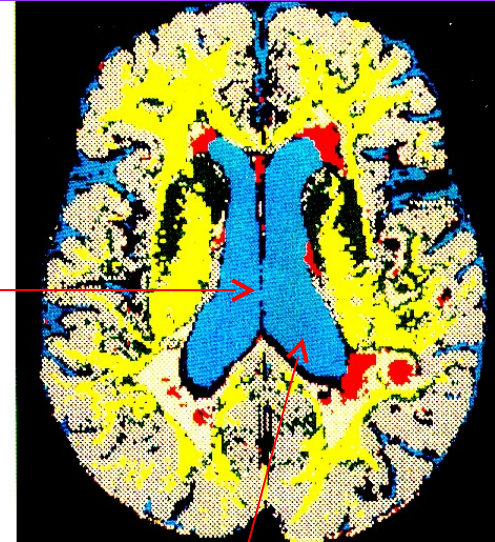
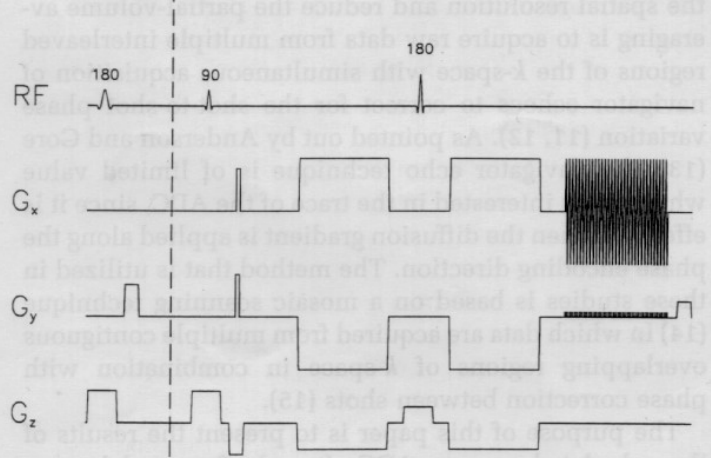
X = 0.000 Y = +0.0000e+0

***** VOXEL NUMBER 595 *****

| Area | ppm | I | Width | |
|---|-------|----------|--------|-----|
| 1st order phasing angle: 50, Pivot point: 4.760 | | | | |
| 2.44e+06 | 3.87 | 2.2e+07 | 0.111 | |
| 1.56e+06 | 3.7 | 4.16e+07 | 0.0376 | |
| 3.3e+06 | 3.61 | 6.96e+07 | 0.0473 | Ins |
| 2.3e+05 | 3.48 | 1.2e+07 | 0.0192 | |
| 6.01e+05 | 3.41 | 2.24e+07 | 0.0268 | |
| 3.55e+06 | 3.26 | 1.33e+08 | 0.0267 | Cho |
| 6.23e+06 | 3.06 | 1.48e+08 | 0.0425 | Cre |
| 1.74e+05 | 2.9 | 1.11e+07 | 0.0158 | |
| 1.52e+05 | 2.84 | 7.58e+06 | 0.02 | |
| 5.15e+05 | 2.69 | 2.68e+07 | 0.0192 | |
| 1.09e+06 | 2.62 | 3.06e+07 | 0.0357 | |
| 1.41e+05 | 2.55 | 5.93e+06 | 0.0238 | |
| 6.36e+05 | 2.47 | 1.98e+07 | 0.0322 | |
| 8.32e+05 | 2.36 | 1.8e+07 | 0.0461 | |
| 1.08e+06 | 2.31 | 2.18e+07 | 0.0494 | |
| 1.91e+06 | 2.16 | 2.87e+07 | 0.0665 | |
| 1.17e+07 | 2.02 | 1.88e+08 | 0.0621 | NAA |
| 3.32e+05 | 1.77 | 8.6e+06 | 0.0386 | |
| 4.22e+06 | 1.53 | 6.11e+07 | 0.0891 | |
| 1.9e+06 | 1.48 | 3.91e+07 | 0.0485 | |
| 1.51e+06 | 1.37 | 2.62e+07 | 0.0575 | |
| 5.88e+05 | 1.24 | 1.57e+07 | 0.0375 | |
| 5.56e+04 | 1.11 | 5.54e+06 | 0.01 | |
| 5.49e+05 | 0.975 | 1.78e+07 | 0.0308 | |
| 2.13e+06 | 0.862 | 2.97e+07 | 0.0718 | |

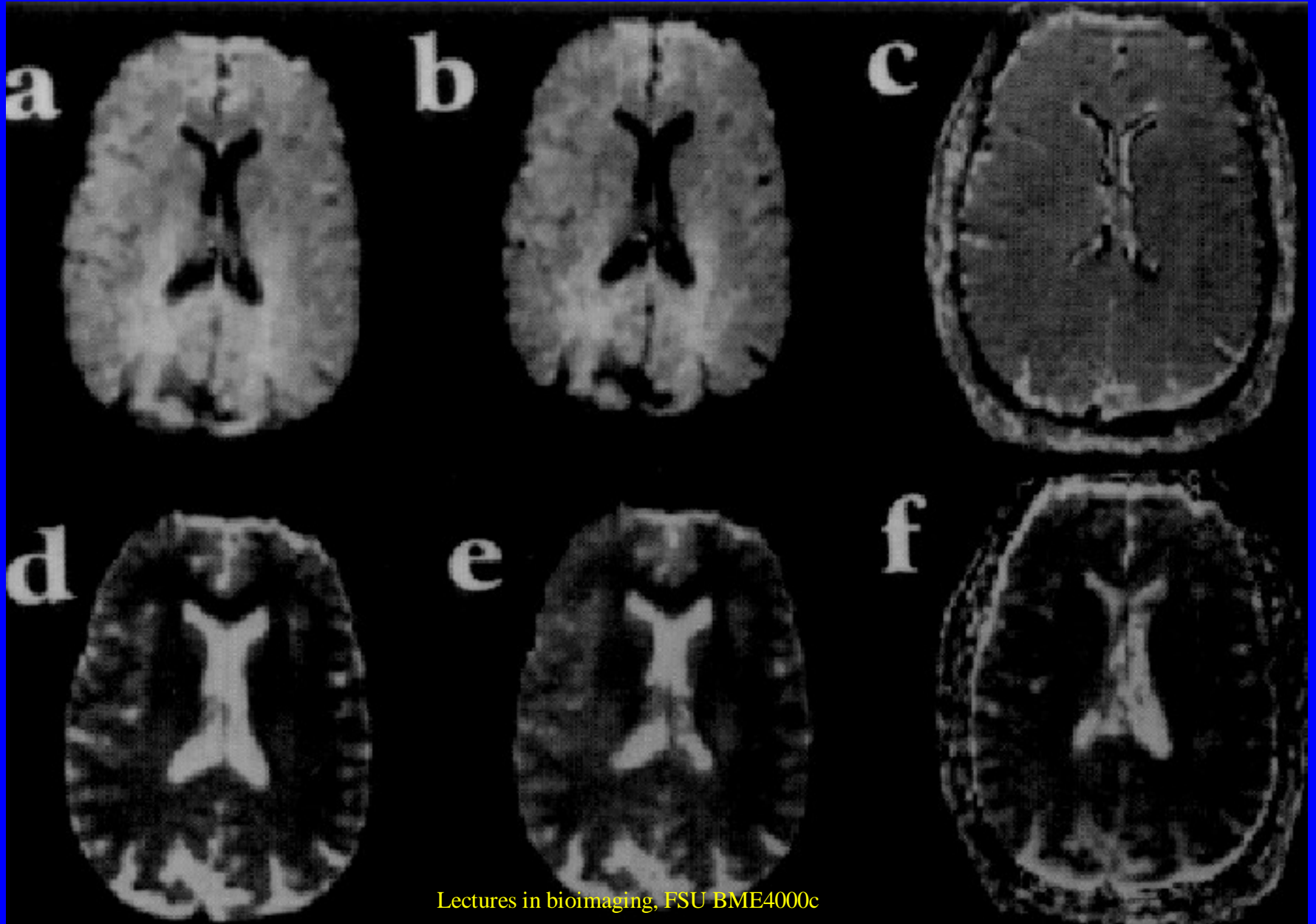
Fast Imaging Technique: EPI Sequence and segmentation

10.1038/npre.2010.4317.1 : Posted 28 Mar 2010



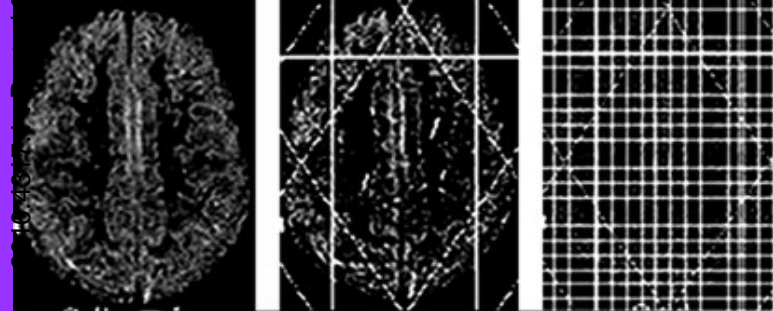
- Segmented color coded features of brain in different regions (MS lesions)

Echo Planar MR Images

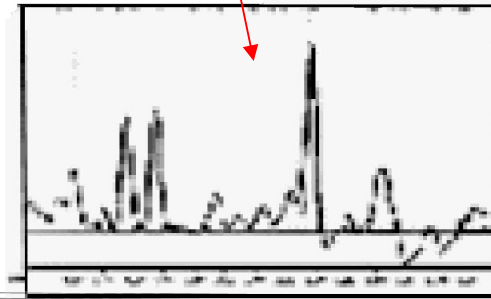
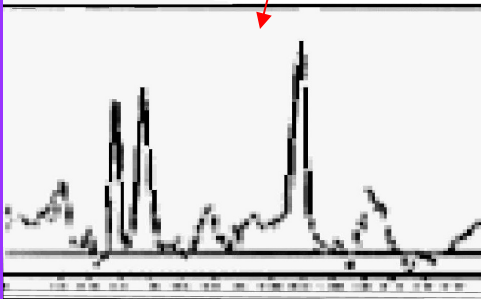
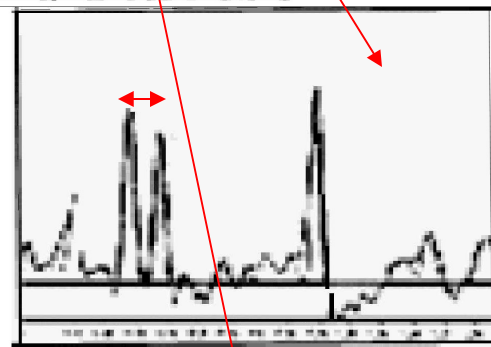
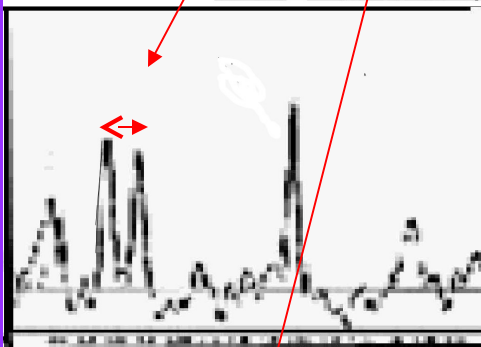
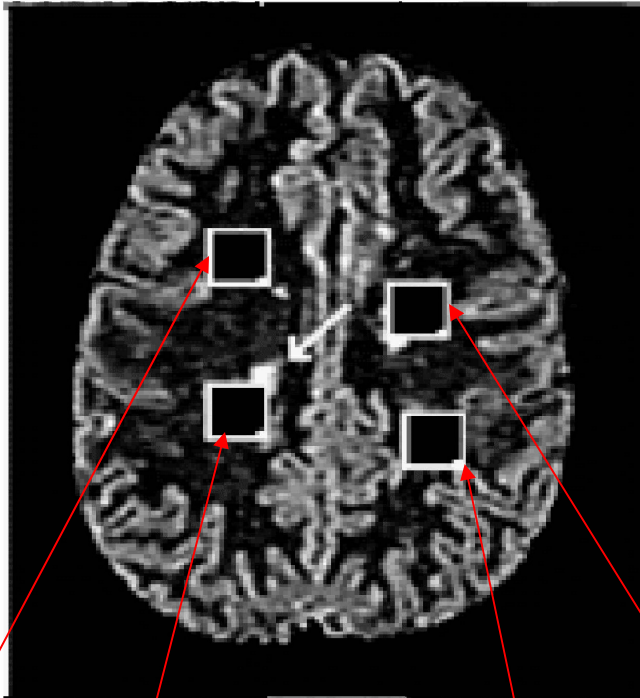
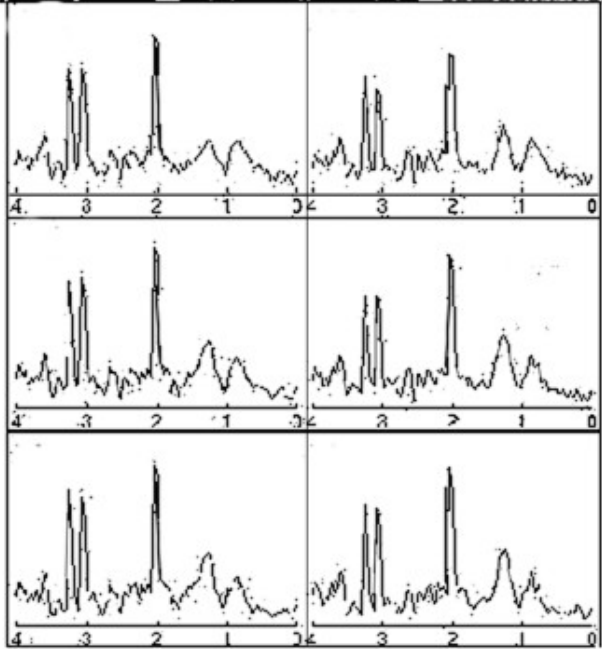


MRSI: MS Lesions High Choline

28 Mar 2010



Nature Precedings : doi:10.1038/n



New Information on MS lesions

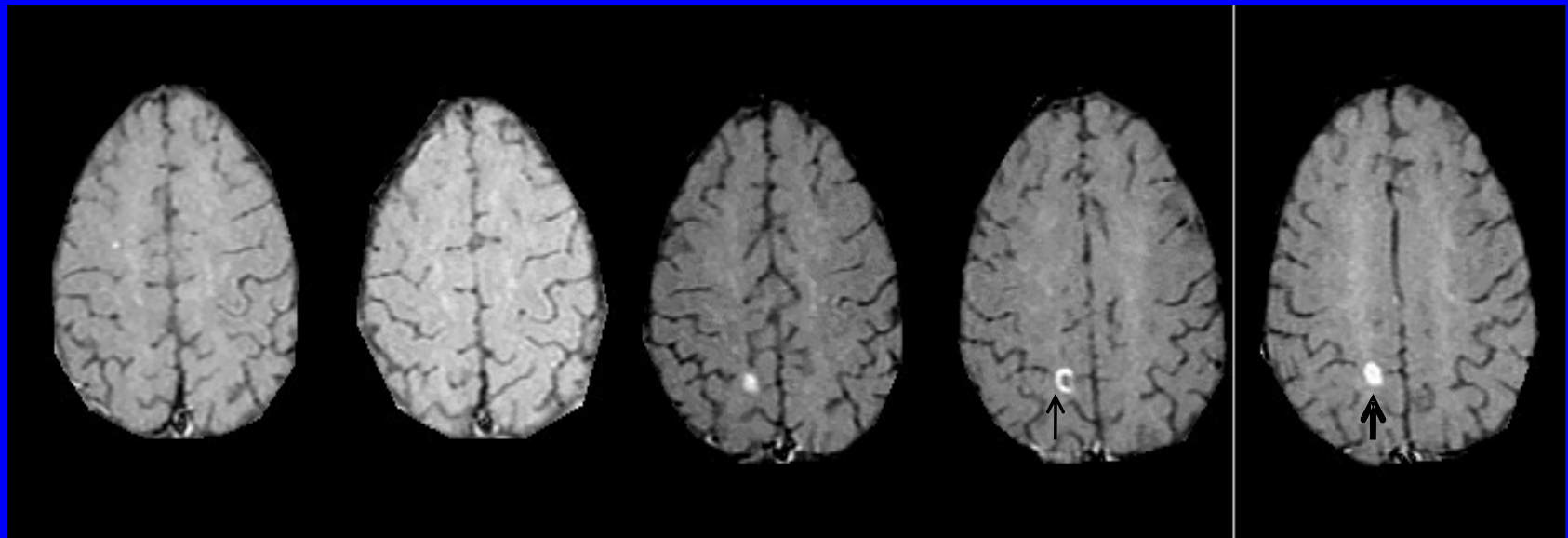
METHODS

- **False Lesion minimization by AFFIRMATIVE**
- **Gray Matter lesions**
- **CSF suppression by Quadruple Contrast sequences**
- **Single 15 mm MRSI slice**
- **32 x 32 Phase encoding steps**
- **Spectroscopic voxel size 0.8 cc; TR=1000 ms, TE=30 ms**

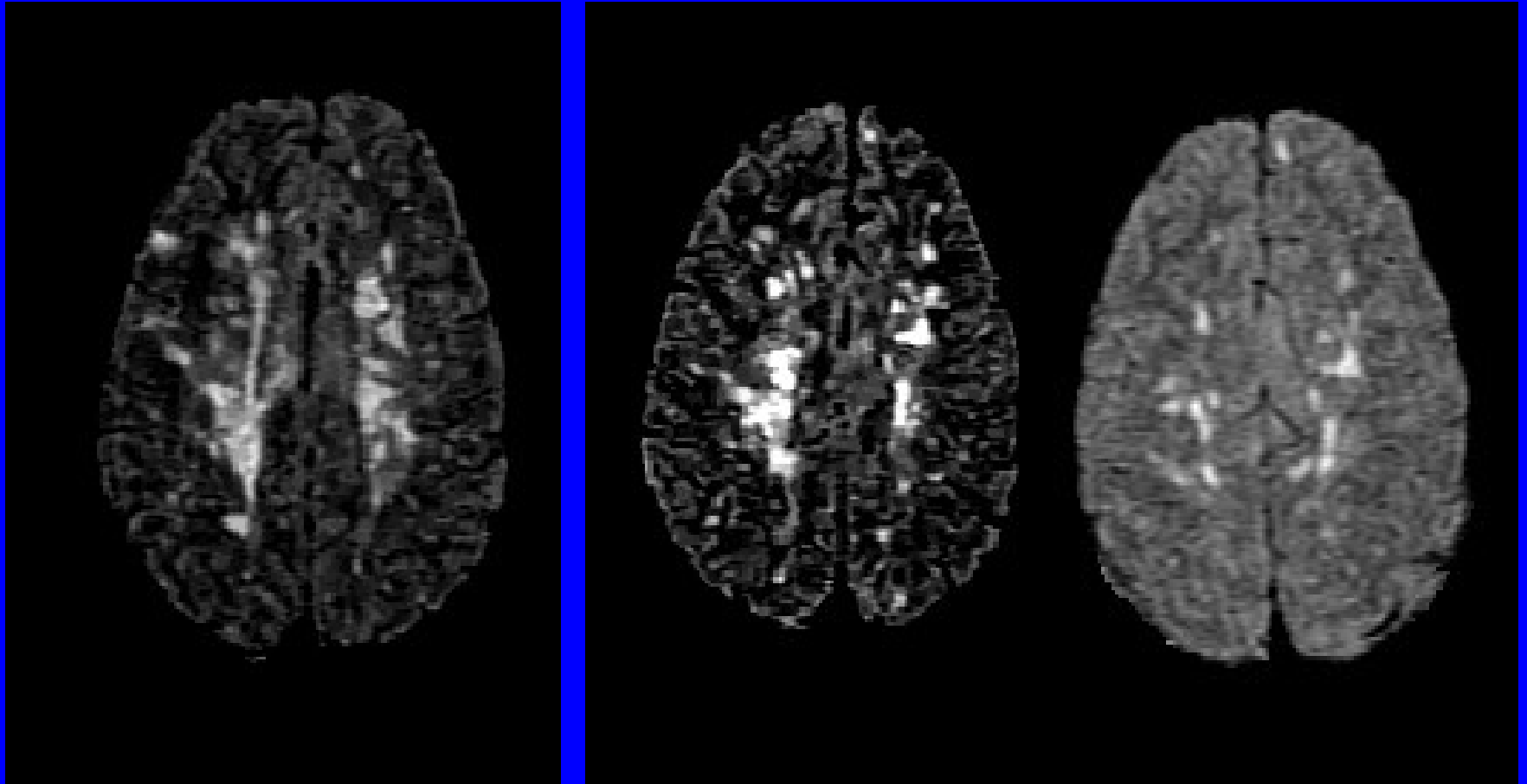
Methods

- **53 Relapsing remitting MS patients examined 1.5 T H-1 MRI and MRSI**
- **3 mm thick axial interleaved contiguous images**
- **MS lesion visualization by AFFIRMATIVE and Quadruple Contrast pulse sequences**
- **MRSI data analysis by APSIP program**

MS lesions showing abnormal appearance in GM and NAGM after Post-Gd Contrast



Localized Acute MS lesion in GM: Pre-Contrast and Post-Contrast images



New Generation of MRI Scanners

- High Magnetic Field Clinical Scanners at 4T, 7T, 11.7T and 17T Strengths

<http://precedings.nature.com/documents/3485/version>

ACKNOWLEDGEMENTS

Proton Magnetic Resonance Spectroscopic Imaging of Abnormal Brain in Multiple Sclerosis

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