4 Tesla 1H MR Spectroscopy and 23Na Imaging in Low Grade Glioma

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Introduction: The potential for high field short echo-time ¹H MR spectroscopy (MRS) to discriminate cell types and determine tumor malignancy from metabolic characteristics continues to grow as the sophistication and reliability of MRS techniques matures. Recent advances in the measurement and removal of the macromolecule baseline from the spectrum further increases the reliability of these measures and allows accurate quantification of glutamate (Glu) and myo-inositol (Myo) in addition to *N*-acetylaspartate (NAA), choline containing compounds (Cho), and total creatine (Cr) (1). Metabolic changes in cerebral tumors are also accompanied by altered tissue sodium concentration reflecting modified cell packing and ion concentrations. The purpose of this study was to characterize the metabolic profile associated with human low-grade glioma and to correlate metabolite levels with changes in tissue sodium signal measured non-invasively by 2^{3} Na MRI.

Methods: Prior to surgery, eight patients (6 male/2 female, mean age 45 ± 11 years) with suspected low-grade glioma were studied using a Varian (Palo Alto, CA) 4T whole body MRI including a Siemens (Erlangen, Germany) Sonata gradient coil. 3D inversion-prepared axial T₁-weighted anatomical images (FOV = 24x24x16 cm³, TI/TR/TE = 500/10/5 ms, 256x256x64 acquisition matrix) were acquired to position a LASER (2) localized single voxel for short echo-time ¹H MRS (128 averages, TR/TE = 2200/46 ms (full spectrum), TI₁/TI₂/TR/TE = 2200/700/4200/46 ms (macromolecule spectrum), 128 averages) within the tumor (biopsy confirmed low grade glioma: N=6, high grade: N=2) and in normal tissue on the contralateral side (N=7). Spectra were lineshape corrected by QUECC (3), residual water subtracted, and the macromolecules baseline removed (1) prior to time domain fitting in fitMAN (1) incorporating prior knowledge of 19 metabolite lineshapes. Metabolite levels were normalized to the unsuppressed water signal from the same voxel after correcting for relaxation effects and water concentration (4) to yield metabolite levels in absolute units (1). 3D axial T₂*-weighted ²³Na images were also acquired (FOV = $24 \times 24 \times 16$ cm³, TR/TE = 23/3.8 ms, 64x64x16 acquisition matrix) and used to calculate sodium signal intensity within each voxel studied by ¹H MRS normalized to the signal intensity in normal parietal-occipital white matter after correcting for T₂ relaxation (5). Metabolite levels were compared (two-tailed ANOVA with p<0.05 considered statistically significant) between low and high-grade tumor and normal tissue and correlated with ²³Na signal intensity.



<u>Results:</u> Figure 1 depicts a typical transverse T_2^* -weighted ²³Na image from one subject with low-grade glioma showing voxel positions (A-tumor, B-control) that correspond to the processed spectra in Figure 2. The tumor is clearly visible in the ²³Na image as a hyperintense region in the right hemisphere. Significantly lower NAA (p<0.01) and Glu (p<0.05, Fig 3) were observed in low and high grade glioma compared to control, while increased levels of Myo (p<0.05 in low grade only), and ²³Na (p<0.01) were also observed. Pooling all data from control and tumor, both Glu and Cr were positively correlated with NAA levels but had significantly different intercepts (p<0.01, Fig p<0.05).

4). ²³Na signal intensity was inversely correlated with both NAA ($r^2=0.59$, p<0.01) and Cr ($r^2=0.37$, p<0.05).

Discussion: Consistent with reduced neuronal density accompanying glial-cell proliferation and previous reports (4), decreased NAA was observed in low-grade glioma. Additional metabolite changes included decreased glutamate and increased myo-inositol. Correlations suggest that all NMR visible glutamate is localized with NAA and is therefore contained within neurons, while the concentration of Cr in glioma cells is ~ 5 mM/L VOI. Increased 23 Na signal in tumors is also consistent with previous studies (6).



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