

In vivo measurement of glycine with short echo-time ^1H MRS in human brain at 7 T

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Abstract

Object To determine whether glycine can be measured at 7 T in human brain with ^1H magnetic resonance spectroscopy (MRS).

Materials and methods The glycine singlet is overlapped by the larger signal of *myo*-inositol. Density matrix simulations were performed to determine the TE at which the *myo*-inositol signal was reduced the most, following a single spin-echo excitation. ^1H MRS was performed on an actively shielded 7 T scanner, in five healthy volunteers.

Results At the TE of 30 ms, the *myo*-inositol signal intensity was substantially reduced. Quantification using LCMoDel yielded a glycine-to-creatine ratio of 0.14 ± 0.01 , with a Cramér–Rao lower bound (CRLB) of $7 \pm 1\%$. Furthermore, quantification of metabolites other than glycine was possible as well, with a CRLB mostly below 10%.

Conclusion It is possible to detect glycine at 7 T in human brain, at the short TE of 30 ms with a single spin-echo excitation scheme.

Keywords Glycine · Magnetic resonance spectroscopy · 7 T · Human brain

Introduction

Glycine is an inhibitory neurotransmitter at glycinergic synapses and an excitatory neurotransmitter at glutamatergic synapses in the adult mammalian central nervous system [1]. It has been shown that glycine administration to schizophrenia patients improves the NMDA receptor function [2]. Furthermore, elevated levels of glycine have been observed in brain tumors and glycine possibly represents a marker for tumor malignancy [3,4]. Thus, it is of great interest to explore non-invasive methods to measure brain glycine.

In general, glycine detection requires specific ^1H magnetic resonance spectroscopy (MRS) editing approaches to overcome the problem of overlap with the much larger resonances of *myo*-inositol. A number of methods have been proposed to this aim: at 3 T, the 2D J-PRESS sequence [5] and another method, which uses multiple refocusing pulses at a very long echo time (TE) [6], and at 4 T, the TE-averaged PRESS sequence [7]. These approaches, however, involve data acquisition at long TEs. With increasing field strength, the Hahn T₂ of metabolites decreases due to an increased dynamic dephasing contribution. The T₂ of NAA, for instance, decreases from ~ 240 ms at 4 T to ~ 160 ms at 7 T [8]. Thus, at high fields, these approaches might suffer from a substantial signal loss, due to data acquisition at long TE.

As the number of MR scanners operating at 7 T is steadily increasing, it is of interest to investigate the potential of MRS at 7 T for in vivo detection of less well represented metabolites, such as glycine, which can have a considerable impact in clinically relevant topics. At 7 T, the high signal-to-noise ratio (SNR) and increased spectral dispersion are expected

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to result in a higher precision of metabolite quantitation and this could allow for investigations of smaller voxel sizes, which better match a specific brain anatomy. However, spectral overlap with the larger resonances of the *myo*-inositol still hampers the detection of glycine and previous MRS studies performed at 7 T reported no detection of glycine [9, 10].

The aim of the present study was to determine whether glycine can be measured at 7 T with a short-TE approach, so as to also allow for quantitation of metabolites other than glycine.

Materials and methods

Theory

The MR detectable protons of glycine generate a singlet at 3.55 ppm, which overlaps with the strongly coupled resonances of *myo*-inositol at 3.52 ppm. As the TE increases, the signal intensity of the glycine singlet decreases due to T2 relaxation. In contrast, the evolution of the coupled resonances of *myo*-inositol is more complex. In addition to the T2 relaxation signal loss, *myo*-inositol displays a J-induced signal dephasing already at a relatively short TE, because of its large J-coupling constants. Consequently, a possible approach for measuring glycine is to identify a TE at which the *myo*-inositol signal intensity is reduced relative to that of glycine. *Myo*-inositol can be modelled as an AM₂N₂P spin system, where the relevant resonances for glycine detection are the M₂ protons at 3.52 ppm and the N₂ protons at 3.61 ppm [11]. In the present study, the spectrum of *myo*-inositol was simulated as a function of TE, for a single spin-echo coherence generation, using the density matrix formalism [12, 13]. Radio-frequency (RF) pulses were modeled as infinitely short, non-selective pulses [12, 13].

MR experiments

All experiments were conducted according to the procedure approved by the institutional review board. Experiments were performed on an actively shielded 7T/68 cm MR scanner (Siemens Medical Solutions, Erlangen, Germany) with a head gradient insert (41 cm, 80 mT/m maximum gradient strength). A home-built quadrature surface coil with two 13 cm-diameter loops was used as a transmitter/receiver. Gradient-echo multislice images were acquired for anatomical localization, to position a 2 × 2 × 2 cm³ voxel of interest (VOI) in the occipital lobe. After first- and second-order shimming with an EPI version of FASTMAP [14], a typical water linewidth in the VOI of interest was 11–12 Hz. Spectra were acquired using the *spin echo full intensity acquired localization* (SPECIAL) scheme, as described more in detail

elsewhere [15]. Briefly, SPECIAL consisted of a spin-echo sequence with a slice-selective 90° asymmetric pulse [16] of 1.28 ms duration and 5.3 kHz bandwidth and a slice-selective 180° pulse [17] of 3.2 ms duration and 1.8 kHz bandwidth. Signal localization in the third dimension was achieved by a 1D ISIS approach, with an adiabatic hyperbolic-secant inversion pulse of 5 ms duration. Prior to signal excitation, VAPOR water signal suppression and outer volume saturation (OVS) were performed [16]. The volume selection performed with a 1D ISIS approach can be more susceptible to subject motion, compared to standard single-shot acquisition approaches such as PRESS and STEAM. On the other hand, the OVS bands applied prior to excitation highly reduce the sensitivity to motion by nulling the signal from tissues surrounding the VOI. As a result, no data degradation due to motion was observed in this study. Spectra at the TE of 30 ms (TR = 4 s, number of averages = 64, spectral width = 4 kHz, 512 ms acquisition time) were acquired in five healthy volunteers. Metabolite concentrations and Cramér–Rao lower bounds (CRLB) were determined by LCModel [18], using a basis set of 21 metabolites.

Results

Density matrix simulations showed that at a very short TE, *myo*-inositol resonances displayed only a small dephasing induced by J-modulation. For TEs up to ~10 ms, the effect of the J-modulation on *myo*-inositol spectral shape was negligible, under single spin-echo excitation (Fig. 1, left). As TE increased, the signal dephasing became more pronounced, in particular for TE in the 30–40 ms range (Fig. 1, right). At the TE of 30 ms, the spectral lineshape of *myo*-inositol reso-

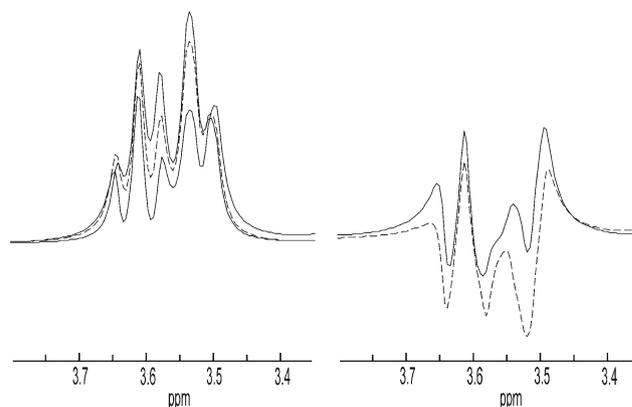


Fig. 1 Density matrix simulations of the M₂ and N₂ *myo*-inositol resonances under single spin-echo excitation at 7 T. *Left* From top to bottom: simulated spectral lineshape at the TE of 2 ms (solid line), 10 ms (dashed line) and 20 ms (solid line). *Right* Simulated spectral lineshape at the TE of 30 ms (solid line) and 40 ms (dashed line)

nances was largely reduced at the resonance frequency of the glycine singlet. Thus, this TE was chosen for in vivo measurements. In vivo ^1H MR spectra at TE = 30 ms showed a substantial signal modulation in *myo*-inositol, as well as in other strongly coupled spin systems, such as the AB spin system of the NAA aspartate moiety at 2.48 and 2.67 ppm, for instance (Fig. 2). When in vivo spectra were fitted to a basis set, which did not include the glycine peak, a large fit residual with positive amplitude was observed at 3.55 ppm (Fig. 3, left). This residual was minimized to the noise level when including glycine in the basis set (Fig. 3, right). Quantification using LCModel yielded a glycine-to-creatine ratio of 0.14 ± 0.01 , with an average CRLB of $7 \pm 1\%$. Furthermore, quantitation of metabolites other than glycine was possible as well, with CRLB mostly below 10% (Table 1). The detection of less well represented metabolites suffered from the signal loss due to the J-evolution of coupled resonances. It should be noted that, despite its lower concentration, the CRLB of glutathione was much lower than that of glutamine (Table 1). This can be ascribed to the fact that the spin system of glutathione includes a singlet at 3.77 ppm, the ‘glycine moiety’ [11], which does not experience signal loss due to J-modulation.

Discussion

In the current study, we show that it is possible to reliably measure glycine at 7 T, at the short TE of 30 ms. Furthermore, at this TE a large number of metabolites can be detected with a CRLB mostly below 10%. The current approach has the advantage of experimental simplicity, since no editing RF pulses or gradient schemes are required to resolve the glycine resonance. In view of the potential application to tumor grading, where multivoxel approaches are necessary, the low CRLB of glycine (<10%) achieved in this study would possibly ensure reliable detection also for voxel sizes smaller than the one used in the current study.

SPECIAL is a novel localization method for in vivo MRS. SPECIAL, like PRESS, is based on a spin-echo acquisition scheme and thus provides full signal intensity from the selected VOI. SPECIAL generates a single spin-echo coherence selection which results in a faster signal dephasing induced by the J-modulation when compared to the double spin-echo coherence selection of PRESS. This is due to the lack of the second refocusing 180° pulse, which in the PRESS sequence partially quenches the signal dephasing induced by the J-modulation. Thus, the advantage of SPECIAL is that a short TE (i.e. 30 ms) can be used, while for PRESS a longer TE would be required to detect glycine. However, at longer TEs, a decrease in SNR is to be expected because of increased T2 signal losses, which are increasingly important at 7 T.

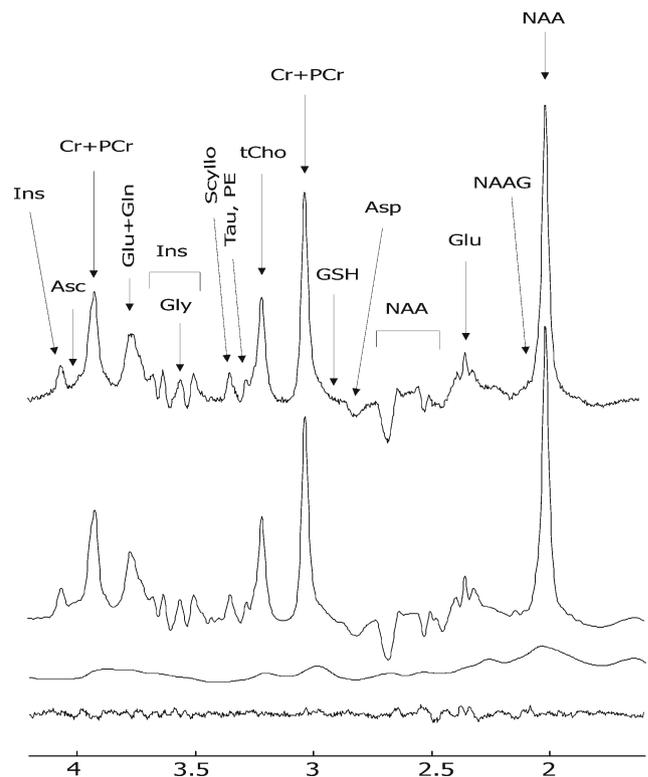


Fig. 2 In vivo ^1H MR spectrum at 7 T, LCModel fit, LCModel fit of the macromolecule spectrum and residuals (from top to bottom), at TE = 30 ms. The spectrum was acquired in a $2 \times 2 \times 2 \text{ cm}^3$ voxel of interest (VOI) in the occipital lobe, with the SPECIAL sequence (*Asc* ascorbate, *Asp* aspartate, *tCho* total choline = GPC (glycerophosphocholine) + PCho (phosphocholine), *Cr + PCr* creatine + phosphocreatine, *Gln* glutamine, *Glu* glutamate, *Gly* glycine, *GSH* glutathione, *Ins* *myo*-inositol, *NAAG* N-Acetylaspartylglutamate, *NAA* N-acetylaspartate, *PE* Phosphoethanolamine, *Scyllo* scyllo-inositol, *Tau* taurine)

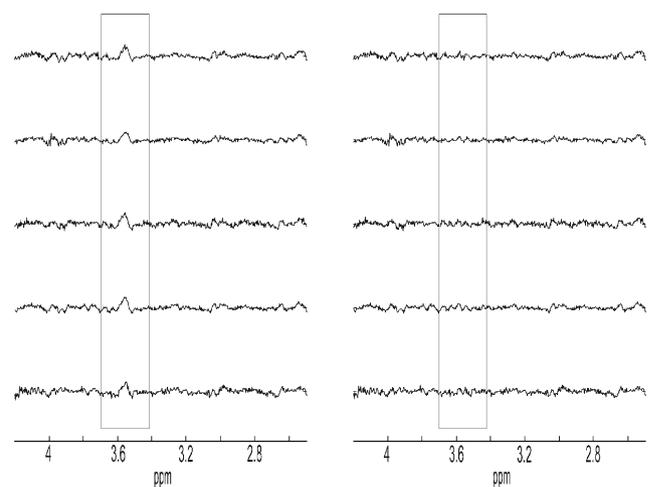


Fig. 3 Residuals obtained from the LCModel fit using a basis set without (left) and with (right) glycine. The box highlights the resonance frequency of the glycine singlet. A large residual with positive amplitude was consistently observed at the glycine resonance frequency in all subjects, when using a basis set without the glycine peak

Table 1 Quantification of metabolite concentration (expressed as a ratio to total creatine, tCr) and Cramér–Rao lower bound (CRLB) at TE = 30 ms

Metabolite	tCr ratio	CRLB (%)
Asc	0.17 ± 0.04	25 ± 9
Asp	0.32 ± 0.03	9 ± 1
Glc	0.28 ± 0.07	17 ± 5
Gln	0.22 ± 0.06	23 ± 12
Glu	1.51 ± 0.13	3 ± 1
Gly	0.14 ± 0.01	7 ± 1
GPC + PCho	0.13 ± 0.01	2 ± 1
GSH	0.18 ± 0.03	7 ± 2
Ins	0.61 ± 0.07	3 ± 1
NAAG	0.12 ± 0.03	10 ± 4
NAA	1.37 ± 0.06	1
PE	0.35 ± 0.03	6 ± 1
Scyllo	0.04 ± 0.01	8 ± 2
Tau	0.41 ± 0.04	10 ± 2

Glc glucose; for the other abbreviations, see Fig. 2 legend

Previously reported values of glycine-to-creatine ratio were in the range of 0.03–1.23, with average values of CRLB in the range of 12–16% [5–7]. The glycine concentration observed in the current study is at the upper limit of the reported values, while the CRLB is substantially smaller. It should be noted that at 7 T the sensitivity of editing methods previously used for glycine detection might suffer from the shortened T₂. With data acquisition at short TE, the quantitation benefits from reduced T₂ losses. Thus, the SNR of the current method is large compared to that of long TE approaches.

Conclusion

In conclusion, glycine can be measured at 7 T in human brain in vivo, at the short TE of 30 ms. In the current approach, no additional RF pulses—other than those needed for signal localization—were required to detect the glycine resonance.

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