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The Ratio of Acetate-to-Glucose Oxidation in Astrocytes from a Single ¹³C NMR Spectrum of Cerebral Cortex

Isaac Marin-Valencia^{1,2}, M. Ali Hooshyar⁷, Kumar Pichumani³, A. Dean Sherry^{3,5,8}, and Craig R. Malloy^{3,4,5,6}

¹ Department of Neurology and Neurotherapeutics, University of Texas Southwestern Medical Center at Dallas, Dallas, TX

² Department of Pediatrics, University of Texas Southwestern Medical Center at Dallas, Dallas, TX

³ Advanced Imaging Research Center, University of Texas Southwestern Medical Center at Dallas, Dallas, TX

⁴ Department of Internal Medicine, University of Texas Southwestern Medical Center at Dallas, Dallas, TX

⁵ Department of Radiology, University of Texas Southwestern Medical Center at Dallas, Dallas, TX

⁶ VA North Texas Health Care System, Dallas, TX

⁷ Department of Mathematical Sciences, University of Texas at Dallas, Richardson, TX

⁸ Department of Chemistry, University of Texas at Dallas, Richardson, TX

Abstract

The ¹³C labeling patterns in glutamate and glutamine from brain tissue are quite different after infusion of a mixture of ¹³C-enriched glucose and acetate. Two processes contribute to this observation, oxidation of acetate by astrocytes but not neurons, and preferential incorporation of α -ketoglutarate into glutamate in neurons, and incorporation of α -ketoglutarate into glutamine in astrocytes. The acetate:glucose ratio, introduced previously for analysis of a single ¹³C NMR spectrum, provides a useful index of acetate and glucose oxidation in the brain tissue. However, quantitation of relative substrate oxidation at the cell compartment level has not been reported. A simple mathematical method is presented to quantify the ratio of acetate to glucose oxidation in astrocytes, based on the standard assumption that neurons do not oxidize acetate. Mice were infused with [1,2-¹³C]acetate and [1,6-¹³C]glucose, and proton decoupled ¹³C NMR spectra of cortex extracts were acquired. A fit of those spectra to the model indicated that ¹³C-labeled acetate and glucose contributed approximately equally to acetyl-CoA (0.96) in astrocytes. Since this method relies on a single ¹³C NMR spectrum, it can be readily applied to multiple physiologic and pathologic conditions.

Correspondence: Craig R. Malloy, MD University of Texas Southwestern Medical Center 5323 Harry Hines Blvd., Dallas, Texas 75390-8813. USA Tel.: 214-645-2720 Craig.Malloy@UTSouthwestern.edu.

INTRODUCTION

Energy-providing substrates are oxidized in the brain at varying rates depending on concentration, nutritional state, age and disease (Schurr 2002, Rothman et al. 1999, Medina & Tabernero 2005, Nehlig & Pereira de Vasconcelos 1993). The biochemical pathways involved in metabolism of each substrate also differ in the two major metabolic compartments. Energy production in neurons is derived primarily from oxidation of glucose (van den Berg & Garfinkel 1971, Gruetter et al. 2001, Oz et al. 2004), whereas oxidation of acetate and glucose as well as net synthesis of glutamate and glutamine occur in astrocytes (Martinez-Hernandez et al. 1977, Waniewski & Martin 1998, Mason et al. 1995). Since astrocytes support neuronal energy production and play an essential role in metabolism of glutamate released from neurons, a major goal has been to quantify biochemical processes in astrocytes separate from neurons. ¹³C NMR spectra acquired over time have been analyzed to measure the rates of oxidation of various ¹³C-labeled energy substrates in both astrocytes and neurons in cerebral cortex (Rothman et al. 1999, Gruetter et al. 2001, Shen 2013, Sibson et al. 2001, Jeffrey et al. 2013, Lanz et al. 2014). Although highly informative, kinetic data are difficult to acquire. A popular alternative has been the analysis of a single highresolution ¹³C NMR spectrum obtained from tissue extracts where it is a simple matter to analyze spin-coupled multiplets.

One consistent finding has been the differences in ¹³C labeling of glutamate and glutamine at carbon 4 (C4) in the presence of acetate. Cerdan and colleagues attributed these differences to preferential oxidation of acetate in astrocytes, qualitatively illustrated in Figure 1 (Cerdan et al. 1990). Differences in ¹³C labeling of glutamate and glutamine have been confirmed in cell preparations, isolated tissues, and in vivo (Deelchand et al. 2009, Hassel et al. 1997, Marin-Valencia et al. 2012, Morris & Bachelard 2003, Haberg et al. 1998, Taylor *et al.* 1996). By presenting a mixture of [1,2-¹³C]acetate and [1-¹³C]glucose, Bachelard, Morris and colleagues (Taylor et al. 1996, Morris & Bachelard 2003) defined the acetate:glucose ratio as $([3,4,5^{-13}C]glutamate + [4,5^{-13}C]glutamate) / ([3,4^{-13}C]glutamate + [4,5^{-13}C]glutamate + [4,5^{-13}C]glutamate$ [4-¹³C]glutamate). In this analysis, labeling in positions 1 and 2 does not significantly influence the ¹³C NMR spectrum at the carbon 4 position of either glutamate (~34.2 ppm) or glutamine (~31.5 ppm). If ¹³C enrichment in position 3 is low, the ratio [4.5-¹³C]glutamate/[4-¹³C]glutamate is also used to assess the acetate:glucose ratio (Haberg et al. 1998, Kondziella et al. 2003). The approach is logical because carbon 2 of acetyl-CoA becomes carbon 4 of α -ketoglutarate in the parent compartment. Consequently, the C4 signal in downstream glutamate or glutamine encodes compartment-specific information about ¹³C labeling in acetyl-CoA. The "acetate:glucose ratio" is a convenient description of the ¹³C spectrum since it is simply the ratio of multiplets in the C4 signal. Although detection of these multiplets is consistent with current knowledge of brain metabolism, it does not provide quantitative information about oxidation of either substrate in either cell compartment.

The purpose of this study was to explore whether information about sources of energy in astrocytes and neurons can be determined from a more detailed analysis of the C4 signal from glutamate and glutamine. A previous mathematical model (Malloy *et al.* 1987, Sherry *et al.* 1992, Malloy*et al.* 1990b, Malloy *et al.* 1988) was extended to accommodate two

compartments based on the widely-accepted assumption that neurons do not oxidize acetate. A sensitivity analysis found that the model accurately returned the ratio of acetate oxidation relative to glucose oxidation in astrocytes. This analysis was tested in studies of normal mouse brain acquired during co-infusion of [1,2-¹³C]acetate plus [1,6-¹³C]glucose. The model provides an approach to determining the ratio of acetate to glucose oxidation in a specific cell compartment from a single ¹³C NMR spectrum.

DESCRIPTION OF THE MODEL

Background

Astrocytes are assumed to oxidize acetate, glucose and other substrates. Neurons are assumed to oxidize glucose and other substrates, but not acetate. In this study, $[1,6^{-13}C]$ glucose and $[1,2^{-13}C]$ acetate were available. With this combination of ¹³C labeling conditions, the possible ¹³C patterns in acetyl-CoA in astrocytes are $[1^2C]$ -, $[2^{-13}C]$ - and $[1,2^{-13}C]$ acetyl-CoA. Consequently, α -ketoglutarate synthesized in astrocytes has only 3 possible labeling patterns in positions 4 and 5: unlabeled, $[4^{-13}C]\alpha$ -ketoglutarate, or $[4,5^{-13}C]\alpha$ -ketoglutarate. By definition, any α -ketoglutarate labeled in position 4 and 5 must have been synthesized in astrocytes. Similarly, α -ketoglutarate synthesized in neurons has only 2 possible labeling patterns in positions 4 and 5: unlabeled or $[4^{-13}C]\alpha$ -ketoglutarate. $[4,5^{-13}C]\alpha$ -ketoglutarate is not generated in neurons.

The covalent bond between position 3 and 4 in α -ketoglutarate is preserved regardless of any subsequent exchanges into glutamate or glutamine. Consequently, for example, if $[4,5^{-13}C]\alpha$ -ketoglutarate or $[3,4,5^{-13}C]\alpha$ -ketoglutarate is generated in astrocytes, the carbon skeleton may exchange into other pools of glutamate and glutamine, as well as the α -ketoglutarate pool of neurons. Regardless of the cell compartment, the labeling pattern in carbons 3, 4 and 5 is preserved and the carbon backbone of $[3,4,5^{-13}C]$ glutamate or $[4,5^{-13}C]$ glutamate arose from condensation of acetyl-CoA with oxaloacetate in astrocytes. The same statement applies to ¹³C labeling in glutamine.

The word "exchange" or a similar term will be used to describe the transfer of a ¹³C labeling pattern from α -ketoglutarate into glutamate or glutamine. The term "acetate:glucose ratio" will refer to the ratio of multiplets observed in the ¹³C NMR spectrum of glutamate (or glutamine) C4 obtained from brain tissue containing both neurons and astrocytes: ([3,4,5-¹³C]glutamate + [4,5-¹³C]glutamate) / ([3,4-¹³C]glutamate + [4-¹³C]glutamate) as defined earlier (Taylor et al. 1996, Morris & Bachelard 2003) (Haberg et al. 1998, Kondziella et al. 2003).

Description of ¹³C Labeling in Acetyl-CoA and in a-ketoglutarate

Quantitative analysis of the ¹³C NMR spectrum of glutamate and glutamine in terms of metabolic pathways requires a description of the ¹³C distribution in acetyl-CoA and in α -ketoglutarate. These variables arise simply from the possible combinations of ¹²C and ¹³C in the acetyl moiety of acetyl-CoA and in α -ketoglutarate, and are independent of any specific model of brain metabolism.

As noted above, three labeling patterns of acetyl-CoA occur in this study: unlabeled acetyl-CoA derived from unlabeled sources, $[2^{-13}C]$ acetyl-CoA derived from $[1,6^{-13}C]$ glucose, or $[1,2^{-13}C]$ acetyl-CoA derived from $[1,2^{-13}C]$ acetate. The following variables describe the ¹³C-labeling pattern in acetyl-CoA in astrocytes:

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fc3a= the fraction of acetyl-CoA in astrocytes that is [1, 2^{-13}Cacetyl-CoA fc2a= the fraction of acetyl-CoA in astrocytes that is [2^{-13}Cacetyl-CoA fc0a= the fraction of acetyl-CoA in astrocytes that is unlabeled
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and in neurons:

 fc_{2n} = the fraction of acetyl–CoA in neurons that is [2-¹³Cacetyl–CoA fc_{0n} = the fraction of acetyl–CoA in neurons that is unlabeled

By definition, $fc0_a + fc2_a + fc3_a = 1$, and $fc0_n + fc2_n = 1$.

A description of ¹³C labeling in α -ketoglutarate in astrocytes and in neurons is also needed. Although there are 32 ¹³C isotopomers of α -ketoglutarate, there are only 16 isotopomers of α -ketoglutarate in astrocytes that have ¹³C in position 4. The variables a – d describe the relative concentrations of these isotopomers of α -ketoglutarate (α KG) in astrocytes, normalized to a sum of 1:

```
 \begin{array}{ll} \mathrm{a}=& [3,4,5^{-13}\mathrm{C}\alpha KG+[1,3,4,5^{-13}\mathrm{C}\alpha KG+[2,3,4,5^{-13}\mathrm{C}\alpha KG+[1,2,3,4,5^{-13}\mathrm{C}\alpha KG\\ \mathrm{b}=& [4,5^{-13}\mathrm{C}\alpha KG+[1,4,5^{-13}\mathrm{C}\alpha KG+[2,4,5^{-13}\mathrm{C}\alpha KG+[1,2,4,5^{-13}\mathrm{C}\alpha KG\\ \mathrm{c}=& [3,4^{-13}\mathrm{C}\alpha KG+[1,3,4^{-13}\mathrm{C}\alpha KG+[2,3,4^{-13}\mathrm{C}\alpha KG+[1,2,3,4^{-13}\mathrm{C}\alpha KG\\ \mathrm{d}=& [4^{-13}\mathrm{C}\alpha KG+[1,4^{-13}\mathrm{C}\alpha KG+[2,4^{-13}\mathrm{C}\alpha KG+[1,2,4^{-13}\mathrm{C}\alpha KG\\ \end{array} \right.
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Assuming that neurons do not oxidize $[1,2^{-13}C]$ acetate, only 8 isotopomers of αKG with ¹³C in carbon 4 are possible in neurons, normalized to a sum of 1:

 $\begin{array}{ll} \mathrm{e}=& [3,4^{-13}\mathrm{C}\alpha KG + [1,3,4^{-13}\mathrm{C}\alpha KG + [2,3,4^{-13}\mathrm{C}\alpha KG + [1,2,3,4^{-13}\mathrm{C}\alpha KG \\ \mathrm{f}=& [4^{-13}\mathrm{C}\alpha KG + [1,4^{-13}\mathrm{C}\alpha KG + [2,4^{-13}\mathrm{C}\alpha KG + [1,2,4^{-13}\mathrm{C}\alpha KG \\ \end{array} \right.$

By definition, a + b + c + d = 1 and e + f = 1.

A Model Relating Metabolic Variables to ¹³C Isotopomers of a-Ketoglutarate

Models of varying sophistication can be used to understand the relationship between ¹³C labeling in acetyl-CoA, metabolic pathways, and the observed ¹³C NMR spectrum of brain tissue. For the purpose of illustrating this analysis, we chose a simple model of brain metabolism, but other models could be used. Anaplerosis was defined as y_a and y_n for astrocytes and neurons, respectively, following earlier conventions (Malloy et al. 1988, Weinman *et al.* 1957, Strisower *et al.* 1952, Malloy *et al.* 1990a). The fraction of anaplerotic substrate in astrocytes that was ¹³C-labeled in position 3 was defined as $fa3_a$ and the fraction of anaplerotic substrate in astrocytes that was unlabeled was defined as $fa0_a + fa3_a = 1$. This variable was provided because under these conditions, [1,6-¹³C]glucose may be

metabolized to $[3^{-13}C]$ pyruvate and subsequently undergo carboxylation to oxaloacetate, an anaplerotic pathway. Since pyruvate carboxylation is considered inactive in neurons, it was assumed that all substrate for anaplerotic reactions in neurons was unenriched and therefore fa0_n = 1. With this assumption, the expressions describing ¹³C enrichment in neurons are relatively simple (Equations 5 and 6). If flux through pyruvate carboxylase in neurons is thought to be significant, then somewhat more complex expressions should be used, analogous to equations 3 and 4. All 6 mathematically – independent metabolic variables of interest (fc3_a, fc2_a, fa3_a, y_a, fc2_n, y_n) should be determined to fully interpret the spectra and describe relative fluxes in both compartments.

The metabolic conditions for this experiment include multiple possible labeling patterns in acetyl-CoA, anaplerosis and metabolism of $[3-^{13}C]$ pyruvate (derived from $[1,6-^{13}C]$ glucose) via pyruvate carboxylation in astrocytes. The relations between the groups of isotopomers, a – f, and the metabolic variables were derived previously (Malloy et al. 1988, Malloy et al. 1990a). For α -ketoglutarate in astrocytes:

$$a = fc3_{a} \cdot (y_{a} \cdot fa3_{a} + fc2_{a} + fc3_{a}) / ((fc2a + fc3_{a}) \cdot (1 + 2y_{a}))$$
[1]

$$b = fc3_{a} \cdot ((1+2y_{a}) - y_{a} \cdot fa3_{a} - fc2_{a} - fc3_{a}) / ((fc2a + fc3_{a}) \cdot (1+2y_{a}))$$
[2]

$$c = fc2_{a} \cdot (y_{a} \cdot fa3_{a} + fc2_{a} + fc3_{a}) / ((fc2a + fc3_{a})^{*} (1 + 2y_{a}))$$
[3]

$$d = fc2_{a} \cdot ((1+2y_{a}) - y_{a} \cdot fa3_{a} - fc2_{a} - fc3_{a}) / ((fc2a+fc3_{a}) \cdot (1+2y_{a}))$$
[4]

For α -ketoglutarate in neurons:

$$e = fc2_n \cdot (fc2_n) / ((fc2_n) \cdot (1+2y_n)$$
 [5]

$$f = fc2_{n} \cdot \left((1+2y_{n}) - fc2_{n} \right) / \left((fc2_{n}) \cdot (1+2y_{n}) \right)$$
[6]

From these relations, it is possible to show that:

$$(a+b)/(c+d) = fc3_a/fc2_a$$
 [7]

Relation of α -Ketoglutarate Isotopomers from Astrocytes and Neurons to the ¹³C NMR Spectrum

The proposed analysis assumes that ¹³C distribution in glutamate may be sensitive to the ¹³C distribution in α -ketoglutarate in either compartment. Consequently, the observed ¹³C isotopomers of glutamate are the sum of glutamate isotopomers arising from exchange with α -ketoglutarate in each compartment. A weighting factor, "g", is defined as the ratio of the ¹³C isotopomers in glutamate from neurons relative to the ¹³C isotopomers in glutamate from astrocytes. The weighting factor, "h", is defined as the ratio of the ¹³C isotopomers in glutamine from neurons relative to astrocytes. These variables, a – h, are

lower case to emphasize that they cannot be observed directly. The next step is to show the relationship of a - h to the experimentally-observed ¹³C NMR spectrum.

The ¹³C NMR spectrum of glutamate detects glutamate in exchange with α-ketoglutarate from astrocytes plus glutamate in exchange with α -ketoglutarate from neurons, weighted by the relative contribution of each compartment. This concept is illustrated in Figure 2. Similarly, the ¹³C NMR spectrum of glutamine detects glutamine in exchange with α ketoglutarate from astrocyte plus glutamine in exchange with α -ketoglutarate from neurons, weighted by the relative contribution of each compartment. The terminology used to describe the carbon-4 resonance was defined previously (Sherry et al. 1992, Malloy et al. 1988, Malloy et al. 1990b). Briefly, the C4 resonance of either metabolite is composed of 4 signals resulting from ${}^{13}C{}^{-13}C$ spin-spin coupling: a singlet (S), a doublet due to J_{34} (D34), a doublet due to J_{45} (D45) and a doublet of doublets or quartet due to J_{34} and J_{45} (C4Q). The relative areas of these multiplets sum to 1. In this notation, the letter "E" refers to multiplets in glutamate and the letter "Q" refers to multiplets in glutamine. After the letter denoting glutamate or glutamine, the abbreviations S, D34, D45 and Q refer to the singlet, doublet due to J_{34} , doublet due to J_{45} and quartet or doublet of doublets, respectively. With these definitions, the multiplets in carbon 4 of the ¹³C NMR spectrum glutamate and glutamine are related to a - h by the following expressions:

$$\begin{split} & \text{E4S} = \left(\text{d} + gf \right) / \left(\text{g} + 1 \right) \quad [8] \\ & \text{E4D34} = \left(\text{c} + ge \right) / \left(\text{g} + 1 \right) \quad [9] \\ & \text{E4D45} = \text{b} / \left(\text{g} + 1 \right) \quad [10] \\ & \text{E4Q} = \text{a} / \left(\text{g} + 1 \right) \quad [11] \\ & \text{Q4S} = \left(\text{d} + hf \right) / \left(\text{h} + 1 \right) \quad [12] \\ & \text{Q4D34} = \left(\text{c} + he \right) / \left(\text{h} + 1 \right) \quad [13] \\ & \text{Q4D45} = \text{b} / \left(\text{h} + 1 \right) \quad [14] \\ & \text{Q4Q} = \text{a} / \left(\text{h} + 1 \right) \quad [15] \end{split}$$

The sum of all multiplets in C4 of glutamate or glutamine are normalized to 1 (E4S + E4D34 + E4D45 + E4Q = 1, and Q4S + Q4D34 + Q4D45 + Q4Q = 1). With these definitions, the acetate:glucose ratio in glutamate = (E4D45 + E4Q)/(E4S + E4D34) = (a + b)/(c+d+g), and the acetate:glucose ratio in glutamine = (Q4D45 + Q4Q)/(Q4S + Q4D34) = (a+b)/(c+d+h). Qualitatively these expressions are simple to interpret and consistent with previous interpretations of the acetate:glucose ratio. For example, if most glutamate

molecules in the sample are in exchange with α -ketoglutarate in neurons (g >> 1), then acetate:glucose ratio in glutamate is small, meaning that glutamate exchanges with α -ketoglutarate in a compartment that does not oxidize acetate, such as neurons. If the contribution of neurons to glutamine is small (h ~ 0), then the ¹³C NMR spectrum of glutamine is dominated by glutamine in exchange with α -ketoglutarate in a compartment that oxidizes acetate, such as astrocytes.

MATERIALS AND METHODS

Animal Preparation, ¹³C Infusions and ¹³C NMR Spectroscopy

The spectra from normal cortex analyzed in this work have been published (Marin-Valencia et al. 2012). Briefly, normal female mice (*Cre*⁻ of CB57BL/6 and FBV/N background, Jackson Laboratory, Bar Harbor) were studied. The right jugular vein was cannulated under general anesthesia. Seven days post-cannulation, mice were loosely confined to a Lucite cage to prevent ambulation. [1,6-¹³C]-glucose and [1,2-¹³C]-acetate were co-administered as a bolus of 0.2 mg/g of body weight for each tracer (in 0.2 mL saline) over 1 min, followed by a continuous infusion of 0.006 mg/g body weight/min for each tracer (in 0.375 mL saline) at 150 μ L/h for 150 min to assure isotopic steady state. Animals were decapitated and the brains rapidly removed (<15 s), followed by extraction of the cerebral cortex for NMR analysis.

Least Squares Fitting and Simulations

The ¹³C NMR spectrum provides 6 mathematically independent measurements. From definitions or prior knowledge, 3 additional relations are known: a + b + c + d = 1, e + f = 1, and d/c = b/a. In addition, the constraint that 1 > a, b, c, d, e, f > 0, and g, h > 0, was included so that positivity conditions for all parameters are realized. In order to determine the parameters a - h that best fit the above equations, we applied the method of nonlinear least square procedure described by Nocedal and Wright (Nocedal J 1999). The sum of the squares of the difference between the actual data and their predicted values based on the chosen parameters is minimized. In order to test the proposed numerical method for finding the model parameters, a set of simulated spectra were prepared (Malloy et al. 1990a), assuming that two independent cell compartments contribute to both glutamate and glutamine. The fractional contribution of ¹³C labeled substrates to acetyl-CoA was allowed to range from 0-1 in each compartment. The fractional contribution of astrocytes to glutamate was allowed to vary from 0-1 with the remaining glutamate provided by neurons. A similar process was applied independently to glutamine. This synthetic data with varying noise was used as input to the computer code to recover the parameters a-h. The influence of natural abundance ¹³C on this analysis is considered in the discussion. The MATLAB code is provided in the supplementary information.

An example of generating simulated spectra from equations 1 - 6 is shown in Table 1. The details are described in the legend. Under these conditions, the observed ¹³C NMR spectrum in carbon 4 of glutamate (E4S, 0.106170; E4D34, 0.780196; E4D45, 0.040584; E4Q, 0.073050) and glutamine (Q4S, 0.127703; Q4D34, 0.362109; Q4D45, 0.182212; Q4Q, 0.327976) are easily calculated. With this set of data from a simulated spectrum,

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metabolic.m returns a=0.4018, b= 0.2232, c = 0.2411, and d = 0.1339. In this example, a large fraction of glutamate reflects the labeling pattern in α -KG of neurons, and a large fraction of glutamine reflects the labeling pattern in α -KG of astrocytes. The ratio of [1,2-¹³C]acetyl-CoA / [2-¹³C]acetyl-CoA in astrocytes is (a+b)/(c+d) = 1.67 = fc3_a/fc2_a. The program accurately recovered the ¹³C acetyl-CoA labeling pattern in astrocytes.

Statistical Analysis

Goodness-of-fit for the fitting program was calculated by finding sum of squares of the residuals. It was less than 10^{-4} for all considered data. Descriptive statistics in Table 2 include mean, standard deviation and % standard deviation calculated in Excel.

RESULTS

¹³C-NMR Spectra from Normal Cortex

The multiplets in the carbon 4 signal of glutamate and glutamine from normal cortex were quite different, as reported earlier (Figure 3) (Cerdan et al. 1990, Deelchand et al. 2009, Hassel et al. 1997, Marin-Valencia et al. 2012, Morris & Bachelard 2003, Haberg et al. 1998, Taylor et al. 1996). Glutamine C4 exhibited a large doublet D45, indicating that a significant fraction of glutamine exchanged with α -ketoglutarate in a compartment capable of oxidizing [1,2-¹³C]acetate. In contrast, glutamate C4 was dominated by the singlet and the doublet D34, suggesting that most of the glutamate exchanged with α -ketoglutarate in a compartment oxidizing [1,6-¹³C]glucose, and that the compartment oxidizing [1,2-¹³C] acetate contributed relatively little to glutamate.

Previously-Defined Acetate: Glucose Ratio Does Not Indicate Unique Metabolic Conditions

To explore the relationship between the ¹³C NMR spectrum and values returned for a – h from fitting the model, spectra were constructed for a wide range of metabolic conditions in both compartments using equations 1 - 6. The contribution of neurons to glutamate and glutamine (g and h) was allowed to vary for each set of generated synthetic data. From these values of a - h, ¹³C NMR spectra of glutamate and glutamine were generated. A subset of results from these simulations is shown in Figure 4. In this set, the ¹³C NMR spectra of glutamate C4 are shown assuming that the fraction of acetyl-CoA derived from glucose is 60% in neurons (the remainder provided by unlabeled substrates such as ketones) and 30% in astrocytes. The fraction of glutamate derived from astrocytes and the fraction of acetyl-CoA derived from acetate in astrocyte were allowed to vary. The previously –defined acetate:glucose ratio was calculated for all conditions illustrated in Figure 4 (Morris & Bachelard 2003). An identical acetate:glucose ratio, 0.42, was observed for the combination of conditions marked with an asterisk in Figure 4. These simulations illustrate that acetate:glucose ratio described earlier does not reflect a unique set of metabolic conditions when two distinct compartments influence the spectrum.

Data from these simulations were also provided to the MATLAB program. As illustrated in Figure 5, the agreement between the actual and recovered $fc3_a/fc2_a$ is optimal as long as astrocyte and neurons contribute somewhat differently to glutamate and glutamine. The percentage error in $fc3_a/fc2_a$ becomes large if g and h are close, within 20%. This result is

not surprising. If g and h are nearly equal, then by definition the multiplets in the ¹³C-NMR spectra of glutamate and glutamine are indistinguishable and the NMR evidence for compartmentation vanishes, exactly as indicated by equations 8 - 15. However, for the case when h differs substantially from g, the simulated result indicates the recovered parameters are reliable and report $fc3_a/fc2_a$.

The signal-to-noise ratio of the ¹³C-NMR spectrum is limited, so it is also important to determine the effect of noise in the spectrum on the variables returned from the least-squares fitting routine. A simple sensitivity analysis was performed to estimate the effects of noise on the accuracy of the parameters returned from the routine. This was accomplished by generating random numbers between -1 and 1, multiplying them by 0.3% of each area of E4S, E4D34, E4D45, E4Q, Q4S, Q4D34, Q4D45 and Q4Q (for Figure 5B) and adding the result to the synthetic data (area for E4S, E4D34, E4D45, E4Q, Q4S, Q4D34, Q4D45 and Q4Q) as input data for calculation of fc3a / fc2a with 0.3% noisy data. The percentage difference between these values and exact values are shown in Figure 5B. Similar procedures were used for finding the effect of 0.6% and 1% noise, shown in Figure 5C and 5D, respectively. Not surprisingly, the analysis is most accurate if signal-to-noise is high. With increased noise, the accuracy of returned values is more limited as illustrated by the scattered distribution of the results. We found with noise of greater than 1%, the recovered fc3a/fc2a is not good even when g and h are quite different.

Relative Rates of Oxidation of ¹³C-Labeled Glucose and Acetate

The data from normal cortex were fit using the MATLAB program in the supplemental information and the results for a – h are summarized in Table 2. The ratio of acetate vs. glucose oxidation in astrocyte was estimated from the fitting results for a – d. The ratio (a +b)/(c+d) (see equation 7) is the ratio of oxidation of $[1,2-^{13}C]$ acetate relative to the oxidation of $[1,6-^{13}C]$ glucose, or 0.96 ± 0.22 (Table 2). These results demonstrate that during infusion of both acetate and glucose, both ^{13}C -labeled substrates are oxidized at approximately equal rates in astrocytes.

Sources of Glutamate and Glutamine

The scaling factor, g, represents the contribution of 13 C – labeled α -ketoglutarate from neurons to total brain glutamate. Similarly, h represents the contribution of 13 C – labeled α ketoglutarate from neurons to total brain glutamine. It should be emphasized that these variables do not directly represent the fraction of glutamate or glutamine derived from neurons because the fraction of α -ketoglutarate that is 13 C labeled cannot be determined from this analysis. Nevertheless, the values of g and h in Table 2 indicate that much of the 13 C labeling in glutamate was derived from of α -ketoglutarate in neurons, and that relatively little of the 13 C labeling in glutamine was derived from of α -ketoglutarate in neurons. These results are consistent with the standard notion that in brain, the labeling pattern in glutamate largely reflects the α -ketoglutarate pool of neurons and that glutamine originates mainly from astrocytes (Martinez-Hernandez et al. 1977, Waniewski & Martin 1998, Mason et al. 1995).

DISCUSSION

The differences in ¹³C labeling in glutamate and glutamine carbon 4 that arise when brain tissue is provided ¹³C-labeled acetate or mixtures of ¹³C-labeled acetate and ¹³C-labeled glucose have been emphasized previously (Cerdan et al. 1990, Deelchand et al. 2009, Hassel et al. 1997, Marin-Valencia et al. 2012, Morris & Bachelard 2003, Haberg et al. 1998, Taylor et al. 1996). Simultaneous infusion of both ¹³C-labeled glucose and ¹³C-labeled acetate is now a popular experimental approach for studies of neurological diseases such as stroke (Pascual *et al.* 1998, Haberg *et al.* 2009), epilepsy (Eloqayli *et al.* 2003, Eloqayli *et al.* 2004), hydrocephalus (Kondziella et al. 2003), diabetic encephalopathy (Garcia-Espinosa *et al.* 2003) and metabolic disorders such as succinic semialdehyde dehydrogenase deficiency (Chowdhury *et al.* 2007). Although there is general agreement that the differences in ¹³C labeling in glutamate and glutamine are due to metabolic compartmentation in brain, there has been no method for compartment-specific quantitation of relative substrate oxidation from a single ¹³C NMR spectrum.

This approach illustrates a number of points. First, although the acetate:glucose ratio defined earlier demonstrates that both acetate and glucose can be oxidized in the brain ADDIN EN.CITE (, there is no direct relation between the previously-defined acetate:glucose ratio and substrate oxidation in a particular cell compartment. As illustrated in Figure 4, dramatically different metabolic conditions may yield the same acetate:glucose ratio. Second, evidence for differences in sources of acetyl-CoA in astrocyte and neurons arises from differences in the ¹³C NMR spectrum of glutamate vs. glutamine. The ¹³C NMR spectrum of glutamate and glutamine do not differ significantly, meaning that g ~ h in the current terminology. Consequently, the ¹³C NMR spectrum of glutamate and glutamate and glutamine may show identical ¹³C-¹³C coupling patterns even if the oxidized substrates in astrocytes and neurons are quite different. Third, the sensitivity of this experimental approach will be greatest if the contribution of the two compartments to glutamate and glutamine differ significantly.

The model also demonstrates the limits on metabolic information that can be derived from the ¹³C NMR spectrum of carbon 4. Even in the very simple model used to test this analysis, there are 6 independent mathematical variables ($fc3_a$, $fc2_a$, $fa3_a$, y_a , $fc2_n$, y_n) that determine the relative concentration of ¹³C isotopomers in the α -ketoglutarate pools of astrocyte and neurons. However, the ¹³C NMR spectrum of glutamate and glutamine C4 provides only 3 mathematically-independent measures of these isotopomers (a, b, and f, since c, d and e are determined by difference from 1 and equation d/c= b/a). Consequently, none of the metabolic parameters specific to each compartment can be determined independently without additional information. However, the ratio of acetate to glucose oxidation can be determined in astrocytes because this ratio is insensitive to the effects of pyruvate carboxylation and ¹³C labeling in pyruvate used in pyruvate carboxylase as shown in equation 7.

Several limitations should be noted. First, if the relative contribution of neurons and astrocyte to ¹³C glutamate and glutamine are the same (equivalent to g = h), then

metabolism in astrocyte and neurons could be quite different, but the difference would be undetectable based on examination of the glutamate and glutamine spectra. Second, signalto-noise must be high, so this analysis is well-suited to analysis of tissue extracts, but likely difficult to apply *in vivo*. Third, in these studies the fractional enrichment of acetate was not determined. Therefore, all results are reported as metabolism of ¹³C-labeled acetate and ¹³C labeled glucose. In the future, these results should be corrected for the measured fractional enrichment of the substrates. This analysis also does not correct for natural abundance ¹³C. Any natural abundance signal will be detected as an excess singlet and hence as a slight over-estimation of the glucose contribution to acetyl-CoA. The analysis assumes that any [1,2-¹³C]acetyl-CoA in brain tissue is derived exclusively from [1,2-¹³C]acetate. Consequently, the analysis does not account for the possibility of pyruvate recycling in neurons under these conditions, which implies significant conversion of oxaloacetate \rightarrow pyruvate \rightarrow acetyl-CoA \rightarrow citrate. Finally, this analysis assumes that steady-state metabolic conditions have been achieved.

In summary, the ratio of acetate to glucose oxidation in astrocytes may be measured from a single ¹³C NMR spectrum. The analysis is optimal for conditions in which the fraction of glutamate and glutamine in exchange with each compartment is substantially different. Current results in normal cortex are consistent with substantial oxidation of acetate relative to glucose in astrocytes, and a dominant contribution of astrocytes to glutamine labeling. The method is applicable to a wide range of conditions.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

ACKNOWLEDGEMENTS

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Abbreviations

aKG	alpha-ketoglutarate
CAC	citric acid cycle
GLN	glutamine
NMR	nuclear magnetic resonance

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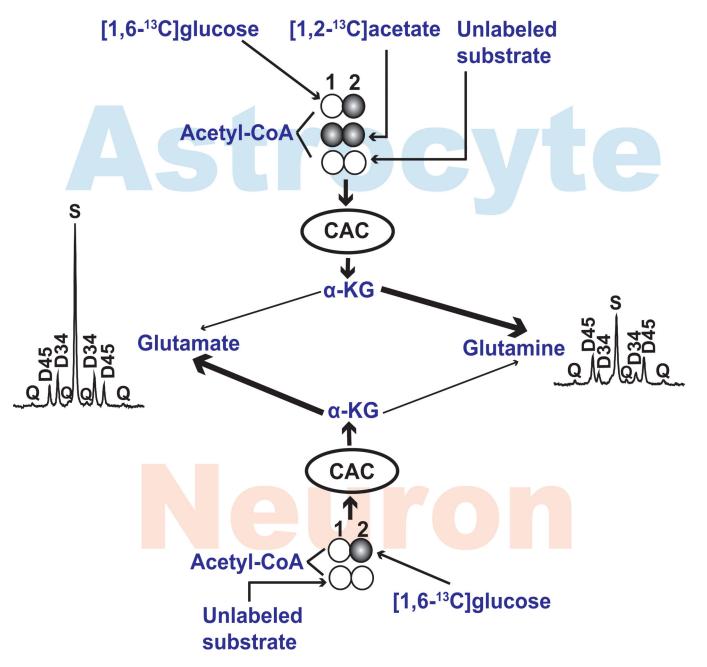


Figure 1. Metabolic Model

The labeling patterns in acetyl-CoA of astrocyte and neurons are illustrated. Acetyl-CoA will be labeled in carbons 1 and 2 from $[1,2-^{13}C]$ acetate, in carbon 2 from $[1,6-^{13}C]$ glucose and it will be unlabeled if derived from unlabeled substrates such as ketones. Acetyl-CoA is oxidized in the citric acid cycle (CAC) to generate α -ketoglutarate. The connecting arrows do not represent detailed metabolic pathways but rather exchange of carbon backbones between α -ketoglutarate and either glutamate or glutamine. The α -ketoglutarate pool in astrocytes can contribute to both glutamate and glutamine; in this example a preferential exchange between α -ketoglutarate in astrocytes with glutamine is illustrated by the heavy line. Similarly, the α -ketoglutarate pool from neurons can exchange with both glutamate and

glutamine; a dominant exchange with glutamate is shown here. Examples of ¹³C NMR spectra from normal mouse cortex illustrate the differences in ¹³C labeling in position 4. Abbreviations: CAC, citric acid cycle; α -KG, α -ketoglutarate; S, singlet; D34, doublet due to J₃₄; D45, doublet due to J₄₅; Q, doublet of doublets or quartet due to J₃₄ and J₄₅. ¹³C-labeled and unlabeled carbons are illustrated with filled circles and unfilled circles, respectively.

GLUTAMATE C4

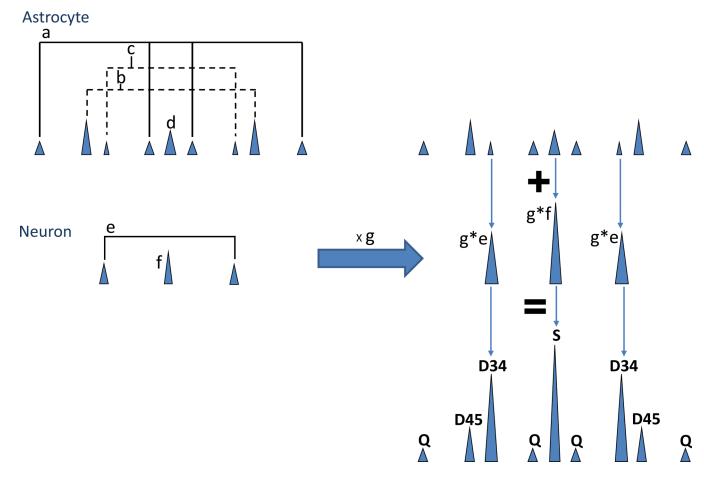


Figure 2. Weighting of the Glutamate Spectrum by Different Contributions from Astrocyte and Neurons

The variables a - g are illustrated. Glutamate derived from astrocyte may contain up to 9 resonances in the its C4 resonance because both [1,2-¹³C]acetate and [1,6-¹³C]glucose may be oxidized in astrocyte. The relative areas of the glutamate labeled in positions 3, 4 and 5 is defined as "a", glutamate labeled in position 4 and 5 as "b", glutamate labeled in position 3 and 4 as "c", and labeled in 4 only as "d". Similarly, the relative areas of glutamate derived from neurons labeled in position 3 and 4 is defined as "e" and the glutamate labeled in position 4 only as "f". The observed glutamate spectrum is the sum of glutamate derived from astrocyte plus the glutamate derived from neurons, weighted by the contribution from neurons, termed "g". A similar analysis applies to interpretation of the glutamine spectrum.

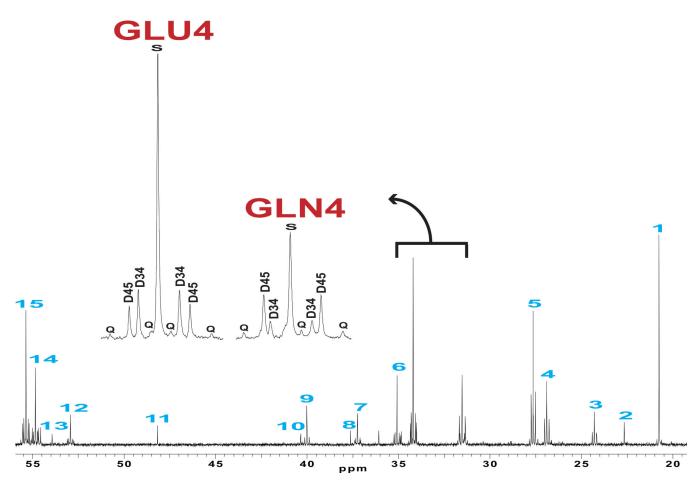
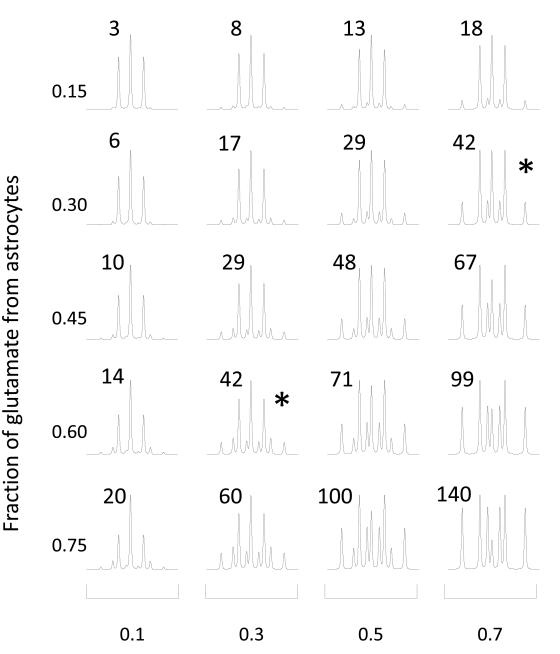


Figure 3. ¹³C NMR spectrum of an Extract from Normal Cortex

The mouse was infused with [1,6-¹³C]glucose and [1,2-¹³C]acetate. The insets display the labeling patterns of glutamate and glutamine in carbon 4. Assignments are: 1, Lactate C3; 2, N-acetylaspartate C6; 3, GABA C3; 4, Glutamine C3; 5, Glutamate C3; 6, GABA C2; 7, aspartate C3; 8, creatine C2; 9, GABA C4; 10, N-acetylaspartate C3; 11, taurine; 12, aspartate 2; 13, N-acetylaspartate C2; 14, Glutamine C2; 15, Glutamate C2; S, singlet; D, doublet; Q, quartet or doublet of doublets; GLN4, glutamine C4; GLU4, glutamate C4.

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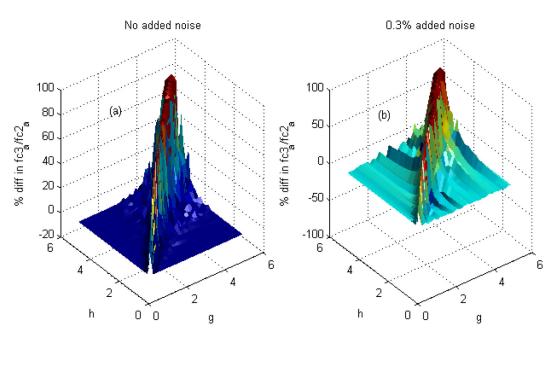


Fraction of acetyl-CoA in astrocytes from [1,2-¹³C]acetate

Figure 4. Simulated ¹³C NMR Spectra of Carbon 4 of Glutamate

These spectra were generated assuming that 60% of acetyl-CoA in neurons is derived from $[1,6^{-13}C]$ glucose and that 30% of acetyl-CoA in astrocyte is derived from $[1,6^{-13}C]$ glucose. Two parameters were allowed to vary: the fraction of glutamate derived from astrocyte (range, 0.15 - 0.75) and the fraction of acetyl-CoA in astrocyte derived from $[1,2^{-13}C]$ acetate (range, 0.1 - 0.7). When astrocytes contribute substantially to glutamate (bottom row) the ¹³C NMR spectrum is very sensitive to changes in metabolism in astrocytes. When astrocytes contribute little to the ¹³C NMR spectrum of glutamate (top

row), the spectrum is insensitive to changes in metabolism in astrocyte. The two spectra marked with "*" had essentially identical acetate:glucose ratios, 42%. The acetate:glucose ratio (in %) for each spectrum is shown in the inset.



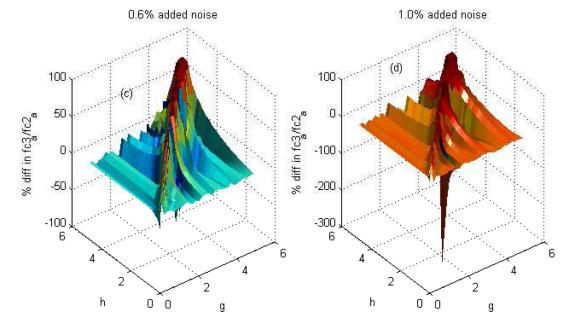


Figure 5. Fitting Simulated Data

Illustrated are the results from analysis of simulated data under different conditions using the program "metabolic.m" in the supplementary material. The procedure is capable of computing a-h most accurately when exchange of glutamate and glutamine with α -ketoglutarate of astrocytes and neurons is quite different. This is demonstrated by finding the difference between the actual ratio of fc2_a (acetyl-CoA labeled in positions 2) and fc3_a (acetyl-CoA labeled in position 1 and 2) for a test model and the estimated fc3_a/fc2_a, using the synthetic data associated with this test model, as the input for the proposed procedure.

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Figure 5A shows that the error is small if g and h are considerably different from each other, but the procedure becomes less accurate in estimating a-h if g and h are nearly equal (error is highest on the diagonal). The noise plays a significant role in the precision of the computation (Figures 5B, 5C and 5D). With increasing the noise from 0 to 1 %, the procedure is less accurate in estimating a-h as represented by significantly worsening error.

Table 1

Example. To illustrate the model and test the fitting algorithm, it was assumed that acetyl-CoA in neurons was derived from [1,6-¹³C]glucose (90%) and unlabeled sources (10%); acetyl-CoA in astrocytes was derived from [1,2-¹³C]acetate (50%), [1,6-¹³C]glucose (30%) and unlabeled sources (20%); flux through pyruvate carboxylase in astrocytes was 20% of flux through citrate synthase; and 50% of the pyruvate undergoing carboxylation was derived from [1,6⁻¹³C]glucose. These assumptions correspond to $fc_{2n} = 0.9$, $fc_{3a} = 0.5$, fc_{2a} = 0.3, ya = 0.2, f_{a3} = 0.5, and yn = 0 in equations 1 – 6. All 32 isotopomers of α -ketoglutarate in neurons or astrocytes were calculated; α -ketoglutarate with ¹³C in position 5 but not 4 cannot occur. The remaining possible isotopomers are shown in 6 groups depending on their appearance in the ¹³C NMR spectrum of C4. The sum of all possible α -ketoglutarate labeling patterns in each compartment is 1. A multiplication factor ("Factor") was used to simulate the concentration in arbitrary units of glutamate or glutamine in exchange with each α -ketoglutarate isotopomer. For example, in neurons α -ketoglutarate labeled in position 4 and 5 is 0.81 of the total α -ketoglutarate synthesized in neurons. The total amount of glutamate in exchange with α ketoglutarate in neurons is $0.81 \times 8 = 6.48$. The sum of each isotopomer group of glutamate or glutamine is shown ("Total"). In this example the amount of glutamate isotopomers labeled in position 4 but not 3 or 5 in the brain is 0.93430. This is the sum of glutamate exchanging in astrocytes (0.21430) plus glutamate exchanging in neurons (0.72). The ¹³C NMR spectrum that would occur in position 4 of glutamate or glutamine was calculated.

	a-Kg	glutamate	glutamine
Neurons		Factor: 8	Factor: 1
[¹² C], [1- ¹³ C], [2- ¹³ C], [1,2- ¹³ C]	0.01000	0.08000	0.01000
[3- ¹³ C], [1,3- ¹³ C], [2,3- ¹³ C], [1,2,3- ¹³ C]	0.09000	0.72000	0.09000
[4- ¹³ C], [1,4- ¹³ C], [2,4- ¹³ C], [1,2,4- ¹³ C]	0.09000	0.72000	0.09000
[3,4- ¹³ C], [1,3,4- ¹³ C], [2,3,4- ¹³ C], [1,2,3,4- ¹³ C]	0.81000	6.48000	0.81000
[4,5- ¹³ C], [1,4,5- ¹³ C], [2,4,5- ¹³ C], [1,2,4,5- ¹³ C]	0	0	0
[3,4,5 ⁻¹³ C],[1,3,4,5 ⁻¹³ C],[2,3,4,5 ⁻¹³ C], [1,2,3,4,5 ⁻¹³ C]	0	0	0
Astrocytes		Factor: 2	Factor: 5
[¹² C], [1- ¹³ C], [2- ¹³ C], [1,2- ¹³ C]	0.07142	0.14284	0.35710
[3- ¹³ C], [1,3- ¹³ C], [2,3- ¹³ C], [1,2,3- ¹³ C]	0.12858	0.25716	0.64290
[4- ¹³ C], [1,4- ¹³ C], [2,4- ¹³ C], [1,2,4- ¹³ C]	0.10715	0.21430	0.53575
[3,4- ¹³ C], [1,3,4- ¹³ C], [2,3,4- ¹³ C], [1,2,3,4- ¹³ C]	0.19287	0.38574	0.96435
[4,5- ¹³ C], [1,4,5- ¹³ C], [2,4,5- ¹³ C], [1,2,4,5- ¹³ C]	0.17857	0.35714	0.89285
[3,4,5 ⁻¹³ C],[1,3,4,5 ⁻¹³ C],[2,3,4,5 ⁻¹³ C], [1,2,3,4,5 ⁻¹³ C]	0.32142	0.64284	1.60710
Total			
[¹² C], [1- ¹³ C], [2- ¹³ C], [1,2- ¹³ C]		0.22284	0.36710
[3- ¹³ C], [1,3- ¹³ C], [2,3- ¹³ C], [1,2,3- ¹³ C]		0.97716	0.73290
[4- ¹³ C], [1,4- ¹³ C], [2,4- ¹³ C], [1,2,4- ¹³ C]		0.93430	0.62575
[3,4- ¹³ C], [1,3,4- ¹³ C], [2,3,4- ¹³ C], [1,2,3,4- ¹³ C]		6.86574	1.77435
[4,5- ¹³ C], [1,4,5- ¹³ C], [2,4,5- ¹³ C], [1,2,4,5- ¹³ C]		0.35714	0.89285
[3,4,5- ¹³ C],[1,3,4,5- ¹³ C],[2,3,4,5- ¹³ C], [1,2,3,4,5- ¹³ C]		0.64284	1.60710

	a-Kg	glutamate	glutamine
¹³ C NMR spectrum			
C4S		0.106170	0.127703
C4D34		0.780196	0.362109
C4D45		0.040584	0.182212
C4Q		0.073050	0.327976

Table 2

Results of Fitting Multiplets from C4 of Glutamate and Glutamine. The variables a - h are reported for cortex from 3 animals. The mean, standard deviation (SD) and % standard deviation (%DEV) are included. The ratio of ¹³C-labeled acetate:glucose oxidation in astrocyte, (a+b)/(c+d), is shown in the final column.

Tissue	a	b	c	d	e	f	g	h	(a+b)/(c+d)
Cortex_#1	0.039	0.478	0.036	0.447	0.164	0.836	2.826	0.205	1.070
Cortex_#2	0.068	0.456	0.062	0.414	0.243	0.756	2.523	0.233	1.101
Cortex_#3	0.055	0.361	0.077	0.507	0.195	0.805	2.032	0.151	0.712
Mean	0.054	0.432	0.058	0.456	0.201	0.799	2.460	0.196	0.961
SD	0.014	0.062	0.020	0.047	0.040	0.040	0.400	0.042	0.216
%DEV	27	14	35	10	20	5.	16.	21.	22.