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# Measuring In Vivo Ureagenesis With Stable Isotopes

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# Abstract

Stable isotopes have been an invaluable adjunct to biomedical research for more than 70 years. Indeed, the isotopic approach has revolutionized our understanding of metabolism, revealing it to be an intensely dynamic process characterized by an unending cycle of synthesis and degradation. Isotopic studies have taught us that the urea cycle is intrinsic to such dynamism, since it affords a capacious mechanism by which to eliminate waste nitrogen when rates of protein degradation (or dietary protein intake) are especially high. Isotopes have enabled an appreciation of the degree to which ureagenesis is compromised in patients with urea cycle defects. Indeed, isotopic studies of urea cycle flux correlate well with the severity of cognitive impairment in these patients. Finally, the use of isotopes affords an ideal tool with which to gauge the efficacy of therapeutic interventions to augment residual flux through the cycle.

# Introduction: Stable Isotopes and the Dynamic Nature of Body Constituents

Fundamental to current understanding of biochemistry is the notion that virtually all molecules of life participate in an incessant cycle of synthesis and degradation that concludes only with death. This conceptualization, so familiar to contemporary science, was not fully grasped until the middle of the 20<sup>th</sup> Century, when researchers first used isotopic tracers in an effort to determine the origin and fate of the molecules that constitute the dense skein of human metabolism.

Two developments facilitated this revolution. The first of these was the discovery by Urey of deuterium at Columbia University in 1933 [1]. The second innovation was the application of isotopic tracers to the study of biochemistry, an effort in large measure spearheaded by Rudolf Schoenheimer, who had emigrated from Germany to the United States. In 1935 he utilized deuterium as a tracer with which to scrutinize intermediary metabolism [2]. Working with the American biochemist David Rittenberg, Schonheimer quickly appreciated the potential utility of the isotopic method for characterizing precursor-product relationships and measuring reaction rates:

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The difficulty in following physiological substances in the course of their transportation in the body and their conversion into other substances, accounts for our ignorance with respect to many of the most fundamental questions concerning intermediate metabolism. The solution of the problems will be possible only when direct methods for tracing such substances will be available... As the chemical properties of the various isotopes of an element are almost identical, it is to be expected that the properties of an organic molecule will remain unaltered if one or even several of the atoms were replaced by their isotopes [1].

The discovery of the urea cycle in 1932 by Krebs and Hensleit [3] was a seminal event, not only because it provided a clear schema by which liver converts waste nitrogen into urea, but because it posited a novel structure – a cycle – that mediated a fundamental biochemical process. Indeed, the urea cycle was the first of the major biochemical cycles to be discovered. As such, it offered a critical insight into the nature of metabolic organization, particularly what Schoenheimer termed the "dynamic state of body constituents" [4]. Now it became apparent that the chemistry of life commonly conformed to a cyclical rather than a linear pattern. The biochemist Joseph Fruton nicely captured the conceptual significance of this paradigm shift:

This work marked a new stage in the development of biochemical thought. Not only was an explanation of a biochemical synthesis offered for the first time in terms of chemical reactions identified in the appropriate biological system and not merely inferred by analogy to the known chemical behaviour of the presumed reactants, but also the paper provided a clue to the organisation of metabolic path ways in living cells. This became evident in 1937 with the appearance of the Krebs citric acid cycle, whose conceptual relation to the earlier ornithine cycle was obvious [5].

#### Isotopic Studies of Urea Turnover

A fortuitous development was the discovery of the urea cycle just prior to the availability of <sup>15</sup>N and <sup>13</sup>C as metabolic tracers [6,7]. Among the first applications of the isotopic method was the demonstration [7] with a <sup>13</sup>CO<sub>2</sub> probe that the carbon atom of urea derived from carbon dioxide (actually,  $HCO_3^-$  is the source). The objective of most early isotopic (usually <sup>15</sup>N) studies of urea synthesis was not to measure flux through the urea cycle but to measure rates of total body protein synthesis [8–10]. In the typical experiment a <sup>15</sup>N-labelled amino acid was administered to human subjects and the appearance of the label in urinary <sup>15</sup>N (most of which is comprised of [<sup>15</sup>N]urea) was determined with isotope ratiomass spectrometry [10]. Whole body protein synthesis is then derived from the assumption that isotope retained in the body corresponds to that incorporated *de novo* into body proteins [8].

Technological advances enabled the use of stable isotopes to study flux through the urea cycle in youngsters with inborn errors of metabolism. Most early studies utilized isotope ratio-mass spectrometry as an analytic tool with which to measure isotopic enrichment. This approach had the advantage of exquisite sensitivity with regard to the detection of isotopic label, but it was far less sensitive with respect to the amount of material that is needed for analysis. In addition, isotope ratio-mass spectrometry requires the complete separation of the analyte of interest and the combustion of this analyze to a gaseous form (CO<sub>2</sub> or N<sub>2</sub>) in order to determine the  ${}^{13}C/{}^{12}C$  or  ${}^{15}N/{}^{14}N$  isotopic ratio. The coupling of the mass spectrometer to a gas chromatograph facilitated pediatric investigations, since reproducible measurements now could be obtained even with relatively small samples sizes (often < 100  $\mu$ l of plasma or serum). In addition, the great resolving power of gas chromatography, which

can separate hundreds of metabolites in a single analysis, greatly amplifies the repertoire of compounds now susceptible to study.

A notable example of this approach is the study of Lee et al [11], who measured urea turnover in a group of patients with urea cycle defects, including ornithine transcarbamylase deficiency (10 female/5 male), citrullinemia (5) and argininosuccinic aciduria (2). The experimental protocol involved the continuous infusion of  $[5^{-15}N]$ glutamine and  $[^{18}O]$ urea, which enabled total body turnover of the urea pool and of the amide-N of glutamine. Measurements of blood  $[^{15}N]$ urea defined the rate of nitrogen transfer from 5-N of glutamine to urea. They found that the ratio of  $[5^{-15}N]$ glutamine/ $[^{15}N]$ urea discriminated the control subjects from those with a urea cycle defect, with this parameter being much higher in patients with disease of neonatal onset.

This study involved a constant infusion of isotope over a relatively prolonged (7.5 hr) period. There are advantages to parenteral administration of tracer, which neutralizes the problem of variable absorption from the gastrointestinal tract. The constant infusion method also lends itself to a straightforward kinetic analysis. However, there are unavoidable problems attendant upon this method, which may be difficult to implement in children. A long infusion period is necessary to achieve steady-state in an individual with a urea cycle defect and consequent diminution of urea turnover. In addition, the relatively small vessels of infants and children commonly present a problem with regard to venous access. Children (as well as many adults) have found it difficult to summon the patience to comply with a long period of isotope infusion. The latter difficulty may be especially vexatious in a child (or adult) with a urea cycle defect, in whom impaired cognition and a relatively short attention span are common comorbidities.

A method based upon oral administration of isotope might prove more practicable to pediatric studies. To this end, we devised a protocol [12,13] based upon the oral administration of  $^{15}NH_4Cl$ . We utilized gas chromatography-mass spectrometry to monitor the appearance of label in [ $^{15}N$ ]urea and [ $^{5-15}N$ ]glutamine over a 4-hour time period following the test dose of tracer. Our study cohort for this X-linked disorder included 15 heterozygotes, of whom 6 were ostensibly asymptomatic, 8 were clearly symptomatic, and 1 was a severely affected individual with neonatal onset.

The results are shown in Fig. 1. We found essentially no difference with regard to the formation of  $[^{15}N]$ urea (left panel) when we compared the response of the control group with that of asymptomatic carriers. In contrast, symptomatic heterozygotes converted significantly less  $^{15}N$  from ammonia to urea. A single, severely affected hemizygote was able to produce virtually no  $[^{15}N]$ urea.

Of note is the finding (Fig. 1, right) that the synthesis of  $[5^{-15}N]$ glutamine was greater even in asymptomatic carriers than in the control group. This observation indicates that ostensibly "asymptomatic" carriers are biochemically distinguishable from normal. In a subsequent study [14] we explored this concept in greater detail by performing neuropsychological testing in 19 female heterozygotes for ornithine transcarbamylase deficiency. The carriers displayed normal IQ scores as well as strength in verbal intelligence and reading, but they manifested weakness in fine motor dexterity and lesser impairments in executive skills, nonverbal intelligence and mathematics. Interestingly, we found that the degree of residual urea synthetic capacity, as measured with the <sup>15</sup>NH<sub>4</sub>Cl loading procedure (Fig. 1), often predicted cognitive measures. Thus, the isotopic method underscored the importance of maintaining good metabolic control, even in otherwise "asymptomatic" carriers. There was a significant positive correlation between individual <sup>15</sup>N incorporation into urea and performance in math (p = 0.0013), attention/executive (p = 0.001) and reading (p = 0.032).

Indeed, the isotopic approach was more predictive of cognitive outcome than the allopurinol challenge test [14].

No method is entirely free of disadvantages. Although the  ${}^{15}NH_4Cl$  loading test proved highly discriminative, it did present problems: (a) The flavor of ammonium chloride is obnoxious and provoked gagging in several subjects. (b) It would be ethically impossible to administer an ammonium salt to an individual who already may have hyperammonemia. (c) Ammonia administration in such a clinical setting is experimentally unsound, since the expanded endogenous N pool would unduly dilute the isotopic tracer. (d) Finally, the method requires gas chromatography-mass spectrometry to analyze enrichment in [ ${}^{15}N$ ]urea and [ $5-{}^{15}N$ ]glutamine. As noted above, this technology is exquisitely sensitive with regard to sample size, but it is several orders of magnitude less sensitive with respect to the detection of isotopic enrichment. This becomes a confounding limitation in studying patients with severe urea cycle defects, in whom we anticipate very little incorporation of label into urea.

To this end, we endeavored to develop a method that would measure residual competency of the urea cycle by quantifying the synthesis of  $[^{13}C]$  urea from a  $^{13}C$ -labeled precursor. The tracer we selected was  $[1-^{13}C]$  acetate, the carbon in which is incorporated into urea via the following reaction sequence:

 $[1 - {}^{13}C] - Acetate \rightarrow TCACycle \rightarrow H^{13}CO_3^- \rightarrow {}^{13}C - Carbamylphosphate \rightarrow [{}^{13}C]Urea$ 

In hepatic mitochondria acetate is very quickly oxidized to bicarbonate following condensation with oxaloacetate in the citrate synthetase reaction. Bicarbonate then enters the urea cycle after reacting with ammonium ion and ATP via carbamylphosphate synthetase. The <sup>13</sup>C-carbamylphosphate ultimately becomes [<sup>13</sup>C]urea. We developed a novel method for determining isotopic abundance in [<sup>13</sup>C]urea. Since this approach deploys isotope ratiomass spectrometry as an analytic tool, it has the advantage of being very sensitive, even when confronted with the challenge of detecting the relatively small amounts of [<sup>13</sup>C]urea that a severely affected patient typically produces.

An important indicator of the value of an isotopic method is whether it yields data that is of clinical utility in terms of devising novel therapies to improve the lives of affected patients. Our group has long been interested in N-carbamylglutamate, a synthetic glutamate derivative that appears to enhance flux through the urea cycle by stimulating the activity of carbamylphosphate synthetase (see above). The drug appears to act by mimicking the action of N-acetylglutamate, which is the naturally occurring obligatory effector of this reaction [15]. We previously showed, using <sup>15</sup>NH<sub>4</sub>Cl as tracer, that N-carbamylglutamate would virtually "cure" defective ureagenesis in patients with a congenital deficiency of N-carbamylglutamate synthetase [16].

We tested  $[1-^{13}C]$  acetate as a probe to see if this species would yield a comparable result. As shown in Fig. 2, when the isotope was given to a patient with a congenital deficiency of N-acetylglutamate synthetase, within minutes it was converted to  $^{13}CO_2$  (Fig. 2, panel A), a peak in which was attained by 30 minutes. This was true before (open circles) and after (closed circles) drug administration. Notwithstanding the robust production of  $^{13}CO_2$ , the patient incorporated relatively little label into  $[^{13}C]$  urea. This was reflected in a paucity of  $^{13}C$  either in the fraction of blood urea that was labeled (Fig. 2, panel B) or in the total concentration of blood  $[^{13}C]$  urea (Fig. 2, panel C). The latter parameter represents the product of the fraction of blood urea labeled with  $^{13}C$  above baseline (Fig. 2, panel B) and the total blood urea concentration. It is evident from the data in Fig. 2 that when this patient

received treatment with N-carbamylglutamate for a period of 3 days, there was a nearcomplete restoration of ureagenesis. Indeed, the post-treatment appearance of label in  $[^{13}C]$ urea is essentially indistinguishable from the control values (data not shown).

In a subsequent study [17] we extended this work to assess the efficacy of N-carbamylglutamate to improve metabolism in patients with propionic acidemia. Hyperammonemia is a common complication in this cohort. The precise mechanism is uncertain, but it probably entails compromise of flux through the carbamylphosphate synthase reaction. We therefore administered a short (3-day) course of N-carbamylglutamate to these patients and again utilized [1-<sup>13</sup>C]acetate to quantify rates of urea production. We again were able to demonstrate a marked increase in [<sup>13</sup>C]urea formation consequent to drug treatment.

The  $[1-^{13}C]$  acetate studies involve administration of a single oral dose of isotopic tracer, an approach we deployed in a "before-and-after" paradigm that entails study on 2 separate days, both prior to and following administration of N-carbamylglutamate. This paradigm could not disclose how quickly subjects respond to the medication – assuming that a response might occur within hours – since label in  $[^{13}C]$  urea does not return to baseline at the end of the 4-hour period of study (see Fig. 2). We therefore developed a method that involves infusion of  $H^{13}CO_3^-$  over a 5-hour period with subsequent sampling of blood to detect  ${}^{13}C$  in  ${}^{13}CO_2$  and  $[^{13}C]$  urea. The rationale for this design is that by administering a priming dose and subsequent constant infusion of  $H^{13}CO_3^-$  we rapidly achieve a steady-state in this tracer since the body bicarbonate pool turns over very rapidly. Hepatocytes then draw upon this pool to form  $[{}^{13}C]$  urea:

 $H^{13}CO_3^- + NH_4^+ + ATP \rightarrow {}^{13}C - carbamylphosphate \rightarrow \rightarrow [{}^{13}C]urea$ 

We found in healthy controls [18] that this approach rapidly (< 10 min) achieves a steadystate in <sup>13</sup>CO<sub>2</sub> in expired air (Fig. 3). Label in plasma [<sup>13</sup>C]urea conformed to a linear pattern (Fig. 4). At 90 minutes after the start of the  $H^{13}CO_3^-$  infusion we gave each subject a single oral dose of N-carbamylglutamate (Fig. 4, arrow). We then continued to monitor label in plasma [<sup>13</sup>C]urea. In 5/6 subjects we noted an upward inflection of the line that described the rate of appearance of plasma [<sup>13</sup>C]urea. These data indicate that Ncarbamylglutamate augments ureagenesis even in healthy individuals and that it does so within 1–2 hours of drug administration.

## Summary

No methodology has contributed more to our understanding of the chemistry of life than the deployment of isotopes by researchers during the past 8 decades. These tracers have enabled the delineation of the remarkable network of interlocking pathways that constitute the rich fabric of human metabolism. The isotopic method has allowed investigators to specify complex precursor-product relationships and to quantify accurately rates of flux through intricate (and otherwise inaccessible) biochemical pathways. Tracer methodology has proved essential to current understanding of human nitrogen metabolism and to our appreciation of the urea cycle as an indispensable mechanism for the elimination of ammonia. Today we recognize that essentially all body proteins participate in the turnover of a myriad of metabolically active nitrogen pools and that they do so with great alacrity, in the process giving rise to the production of copious amounts of waste nitrogen. Perhaps the most dramatic proof of this conceptualization derives from our clinical experience with patients who have inherited urea cycle defects, in whom levels of ammonia rapidly rise to potentially neurotoxic and even lethal concentration. Isotopes have allowed us to quantitate

the dimension of such derangements, to identify the sources of ammonia, and to begin to define the mechanisms by which hyperammonemia works its havoc on the nervous system. Finally, with the isotopic method we are in possession of a powerful tool by which to gauge the efficacy of experimental treatments to enhance ureagenesis and, in so doing, to provide a better outcome for the ever growing cohort of patients who suffer with this terribly debilitating group of diseases.

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#### References

- 1. Urey HC. The Separation and Properties of the Isotopes of Hydrogen. Science 1933:566–571. [PubMed: 17801686]
- 2. Schoenheimer R, Rittenberg D. Deuterium as an indicator in the study of intermediary metabolism. J Biol Chem 1935;111:163–168.
- Krebs HA, Henseleit H. Untersuchungen über die Harnstoffbildung im Tierkörper. Zeitschrift f
  ür physiologische Chemie 1932;210:33–46.
- Schoenheimer, R. The Dynamic State of Body Constituents. Harvard University Press; Cambridge, MA: 1942.
- 5. Fruton, JS. Molecules and Life: Historical Essays on the Interplay of Chemistry and Biology. New York: Wiley Interscience; 1972. p. 436
- Schoenheimer R, Rittenberg D, Foster GL, Keston AS, Ratner S. The application of the nitrogen isotope N15 for the study of protein metabolism. Science 1932;88:599–600. [PubMed: 17831794]
- 7. Rittenberg D, Waelsch H. The source of carbon for urea synthesis. J Biol Chem 1940;136:799-800.
- Duggleby SL, Waterlow JC. The end-product method of measuring whole-body protein turnover: a review of published results and a comparison with those obtained by leucine infusion Brit. J Nutr 2005;94:141–153.
- Koletzko B, Demmelmair H, Hartl W, Kindermann A, Koletzko S, Sauerwald T, Szitanyi P. The use of stable isotope techniques for nutritional and metabolic research in paediatrics. Early Hum Devel 1998;53:S77–S97. [PubMed: 10102657]
- San Pietro SA, Rittenberg D. A study of the rate of protein synthesis in humans. J Biol Chem 1953;201:445–455. [PubMed: 13044814]
- Lee B, Yu H, Jahoor F, O'Brien W, Beaudet AL, Reeds PR. In vivo urea cycle flux distinguishes and correlates with phenotypic severity in disorders of the urea cycle. Proc Natl Acad Sci USA 2000;97:8021–8026. [PubMed: 10869432]
- 12. Yudkoff M, Daikhin Y, Nissim I, Jawad A, Wilson J, Batshaw M. In vivo nitrogen metabolism in ornithine transcarbamylase deficiency. J Clin Investigator 1996;98:2167–2173.
- Yudkoff M, Daikhin Y, Wilson JM, Batshaw. In vivo measurement of ureagenesis with stable isotopes. J Inherit Met Dis 1998;21(Suppl 1):21–29.
- Gyato K, Wray J, Huang ZJ, Yudkoff M, Batshaw ML. Metabolic and neuropsychological phenotype in women heterozygous for ornithine transcarbamylase deficiency. Ann Neurol 2004;55:80–86. [PubMed: 14705115]
- Gebhardt B, Vlaho S, Fischer D, Sewell A, Bohles H. N-carbamylglutamate enhances ammonia detoxification in a patient with decompensated methylmalonic aciduria. Mol Genet Metab 2003;79:303–304. [PubMed: 12948747]
- Caldovic L, Morizono H, Daikhin Y, Nissim I, McCarter RJ, Yudkoff M, Tuchman M. Restoration of ureagenesis in N-acetylglutamate synthase deficiency by N-carbamylglutamate. J Pediatr 2004;142:552–554. [PubMed: 15480384]
- 17. Tuchman M, Caldovic L, Daikhin Y, Horyn O, Nissim I, Korson M, Burton B, Yudkoff M. Ncarbamylglutamate markedly enhances ureagenesis in N-acetylglutamate deficiency and propionic

acidemia as measured by isotopic incorporation and blood biomarkers. Pediatr Res 2008;64:213–217. [PubMed: 18414145]

 Ah Mew NA, Payan I, Daikhin Y, Nissim I, Nissim I, Tuchman M, Yudkoff M. Effects of a single dose of N-carbamylglutamate on the rate of ureagenesis. Mol Genet Metab 2009;98:325–330. [PubMed: 19660971]



#### Figure 1.

Left: Appearance of [<sup>15</sup>N]urea in blood after oral administration of <sup>15</sup>NH<sub>4</sub>Cl (~ 0.37 mmol/ kg; 98 atom % excess) to health controls (closed circles); asymptomatic heterozygotes with ornithine transcarbamylase deficiency (open circles); symptomatic heterozygotes (open boxes); or a severely affected hemizygote with neonatal onset disease (open triangles). Data are mean  $\pm$  SEM at each time point. Right: Appearance of label in blood [5-<sup>15</sup>N]glutamine after administration of the <sup>15</sup>NH<sub>4</sub>Cl. Symbols same as those for the left panel. The "absolute" values refer to the product of the blood concentration of either urea or glutamine and the isotopic enrichment (atom % excess) at each time point. Adapted from Yudkoff et al [12].



#### Figure 2.

A patient with N-acetylglutamate synthetase deficiency was given an oral dose of  $[1^{-13}C]$  acetate and samples subsequently were taken at the indicated times for analysis of <sup>13</sup>C enrichment (atom % excess) in breath <sup>13</sup>CO<sub>2</sub> (Panel A), plasma  $[^{13}C]$  urea (Panel B) and the absolute concentration of  $[^{13}C]$  urea in blood (Panel C). The study was performed both before (open circles) and after a 3 day course of treatment with N-carbamylglutamate. From Caldovic et al [16].



#### Figure 3.

Isotopic abundance (atom % excess) in  ${}^{13}$ CO<sub>2</sub> in expired air in healthy adults who received an infusion of NaH<sup>13</sup>CO<sub>3</sub> (0.089 mmol/kg/hr; 98 atom % excess) for a period of 300 minutes. The solid horizontal line denotes the mean isotopic abundance from 10 to 300 minutes. During the initial 10 minutes of the experiment each subject also received a priming dose (0.083 mmol/kg) of the labeled bicarbonate. At 90 minutes each individual received an oral dose of NCG (100 mg/kg).



#### Fig. 4.

The concentration of  $[^{13}C]$  urea in plasma during the constant infusion of NaH<sup>13</sup>CO<sub>3</sub>. The line represents linear regression analysis from 10 to 150 minutes. At 90 minutes each subject received an oral dose (50 mg/kg) of N-carbamylglutamate. The closed circles correspond to the period during which the linear regression was performed. The open circles are points from 150 until 300 minutes and correspond to the post-treatment values. The time from 10 until 150 minutes is included in the regression because this is a period of drug absorption. The number at the top of each graph (µmol/min/liter) correspond to the rate of  $[^{13}C]$  urea appearance in blood.