

Rates of Urea Synthesis in the Human Newborn: Effect of Maternal Diabetes and Small Size for Gestational Age

SATISH C. KALHAN

Department of Pediatrics, Case Western Reserve University, School of Medicine, Cleveland, Ohio 44106

ABSTRACT. The rates of urea synthesis and glucose turnover were quantified during fasting using [$^{15}\text{N}_2$]urea and [6,6 $^2\text{H}_2$]glucose tracers with the prime constant rate infusion technique in 33 infants: 19 normal infants, 10 infants of diabetic mothers (IDM), and four small for gestational age (SGA) infants born at term gestation. All infants were studied during fasting 4 h after their last feed. Eleven normal infants and six IDM were studied soon after birth before any feeding. The rate of urea synthesis in normal infants was 5.84 ± 2.0 mg of nitrogen (N)/kg·h $^{-1}$ or 3.5 μmol of urea/kg·min $^{-1}$. The rate was slightly higher in IDM (7.09 ± 3.0 mg N/kg·h $^{-1}$) and lower in SGA infants (4.59 ± 1.22 mg N/kg·h $^{-1}$); however, the differences were not statistically significant. No differences in urea synthesis were observed between infants studied soon after birth and those studied after initiation of feeding. The rate of appearance of glucose was lower in IDM infants studied during the first 6 h after birth (IDM 19.62 ± 2.14 μmol /kg·min $^{-1}$, normal infants 24.03 ± 4.05 μmol /kg·min $^{-1}$, $p = 0.01$). However, rate of appearance of glucose in IDM infants studied between 17 and 20 h after birth was similar to that in normal infants. Rate of appearance of glucose was lower (not significantly) in SGA infants (17.7 ± 3.3 μmol /kg·min $^{-1}$) as compared with normal infants. No correlation between rates of urea synthesis and glucose turnover was observed. These data show that newborn infants during fasting have an obligatory rate of protein oxidation of ~ 0.87 g/kg·d $^{-1}$ and that maternal diabetes has no effect on it. The slightly lower rate of protein oxidation in SGA infants may be related to increased N assimilation. (*Pediatr Res* 34: 801–804, 1993)

Abbreviations

N, nitrogen
IDM, infants of insulin-dependent diabetic mothers
SGA, small for gestational age
AGA, appropriate for gestational age
Ra, rate of appearance

Measurements of urea N excretion are performed to quantify nitrogen balance and rates of protein oxidation in infants and children. However, limited data are published in literature on the rates of urea synthesis or excretion in the human newborn. Previous data suggested that the rates of urea N excretion in the newborn infant may be much lower than those in the adults as

a consequence of active growth (1). A later study by Jones *et al.* (2) showed that the rates of urea N excretion in the urine of newborn infants were higher than previously suggested. Nevertheless, the urea excretion rate was much less than that estimated for the human fetus at term gestation by Gresham *et al.* (3). Inasmuch as urinary urea does not reflect the excretion of the total urea synthesized because of loss of urea by skin and possibly because of hydrolysis in the gut, the measurements of urea N excretion in urine will underestimate the actual rates of urea synthesis and therefore protein oxidation (4, 5). In fact, the rates of protein oxidation, as measured by oxidation rates of essential amino acids such as leucine (6, 7), suggest a much higher rate of protein oxidation in full-term newborn infants (~ 1.4 g/kg·d $^{-1}$) as compared with those by the urea excretion rate from previous studies (~ 0.4 g/kg·d $^{-1}$).

In the present study, the rates of urea synthesis were quantified by the tracer dilution technique, which will estimate the total rate of urea synthesis and therefore should provide an estimate of total protein oxidation. Normal full-term infants were studied during fasting, and the effect of maternal diabetes and intrauterine growth retardation on urea synthesis was examined. Because ureagenesis is considered a concomitant of gluconeogenesis from amino acids (8, 9), the rate of glucose turnover was also quantified simultaneously by tracer dilution to examine the relation between ureagenesis and glucose turnover.

MATERIALS AND METHODS

Urea and glucose kinetics were measured in 33 newborn infants during the first 3 d after birth. All infants were studied during fasting, either before the initiation of feeding soon after birth or 4 h after their last feeding. Nineteen were full-term, AGA infants (Table 1). Of these, 11 were studied soon after birth during the first 5 h and eight were studied after the feeding was started on the second or third day. They were fed commercial formula 60–70 mL/kg·d $^{-1}$ (Similac, Ross Laboratories, Columbus, OH) or were breast fed. Of the 10 IDM, six were studied during the first 5 h after birth, and four were studied after the feedings were started on the second day. Four SGA infants were studied between 8 and 36 h after birth. All infants were born after an uncomplicated delivery, did not have any perinatal problems such as asphyxia or sepsis, and were cared for in the newborn nursery for normal infants. Written informed consent was obtained from one or both parents after the procedure was fully explained to them. The study protocol was reviewed and approved by the Institutional Committee on Investigations in Humans. The investigator was not responsible for the clinical care of the mothers or infants. All infants were studied in the Perinatal Clinical Research Center at Cleveland Metropolitan General Hospital in a study room maintained at 30°C and relative humidity of 40%. Two i.v. 23-gauge butterfly needles were placed, one each in the dorsum of the hand and the foot, to infuse the tracers and draw blood samples. The sampling site

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Correspondence: Satish Kalhan, M.D., Division of Neonatology, Rainbow Babies & Childrens Hospital, 2101 Adelbert Rd., Cleveland, OH 44106.

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Table 1. Clinical characteristics of study infants ($n = 33$) (mean \pm SD)*

	Birth weight (g)	Gestational age (wk)	Age at study (h)	Males/females	Delivery (vaginal/cesarean section)
Normal infants					
Not fed ($n = 11$)	3164 \pm 367	39 \pm 1.3	3 \pm 1	6/5	6/5
Previously fed ($n = 8$)	3139 \pm 781	40 \pm 1.9	(8-72)†	3/5	5/3
IDM					
Not fed ($n = 6$)	3612 \pm 611‡	38 \pm 1	3 \pm 1	3/3	1/5
Previously fed ($n = 4$)	3403 \pm 668‡	39 \pm 1	(17-20)†	2/2	3/1
SGA infants					
Previously fed ($n = 4$)	2453 \pm 105‡	41 \pm 0.6	(8-33)†	2/2	4/0

* All infants were studied during fasting. "Not fed" infants were studied soon after birth before any feeding. "Previously fed" infants were studied after the initiation of feeding; however, they had also fasted for 4 h before the start of the tracer infusion, which was continued for 4-5 h.

† Range.

‡ Significantly different when compared with normal infants (analysis of variance, $p < 0.01$).

was kept patent by infusion of normal saline at 2 mL/h along with intermittent heparin saline flushes. Plasma glucose was monitored throughout the study to assure that no infant became hypoglycemic.

[$^{15}\text{N}_2$]urea, 99 atom % ^{15}N , and [6,6- $^2\text{H}_2$]glucose, 98 atom % ^2H , were obtained from Merck & Co, Inc., Dorval, Canada. The tracers were prepared for i.v. infusion and were tested for sterility and pyrogenicity as described previously (10). Both tracers were infused simultaneously as a prime-constant rate infusion via an occlusive pump (Holter pump, Extracorporeal Medical Specialties, Inc., King of Prussia, PA). To achieve an early steady state, usually between 2 and 3 h for urea and ~ 90 min for glucose, a priming dose of both tracers was given. For [$^{15}\text{N}_2$]urea, the bolus prime was 2 mg/kg body weight and the rate of constant infusion was 0.2 mg/kg \cdot h $^{-1}$. For [6,6- $^2\text{H}_2$]glucose, the prime equaled 90 min of infusion or 2.7 mg/kg body weight, and the rate of constant infusion was 60 $\mu\text{g}/\text{kg} \cdot \text{min}^{-1}$. The tracers were infused for a period of 4-5 h. Twenty infants (10 normal infants, six IDM, and four SGA infants) also received [^{13}C]lactate for lactate turnover studies (data reported separately). The rate of infusion was checked in each study gravimetrically at the end using the same needle and infusion system. Blood samples (1.0 mL) were drawn at 30-min intervals in heparinized syringes. Sample size was reduced in smaller infants. Plasma was separated immediately by centrifugation at $\approx 4^\circ\text{C}$ and stored at -10°C for later analysis.

Assays. Plasma glucose was measured by a coupled enzymatic assay using hexokinase and glucose-6-phosphate dehydrogenase on an automated centrifugal system (Baker Instruments, Pleasantville, NY). Plasma urea N was measured on the same system by a coupled reaction using urease and glutamic dehydrogenase (10).

Plasma C-peptide levels were measured by a double antibody radioimmunoassay after extraction of the interfering antibodies by polyethylene glycol (11).

The ^2H enrichment of glucose was measured on an HP5985 gas chromatography mass spectrometry system (Hewlett-Packard, Palo Alto, CA) after the preparation of a pentacetate derivative, as described previously (10, 12). Electron impact ionization (70 eV) was used and m/z 98 and 100 were monitored using selected ion monitoring. Standard glucose solutions of known enrichment were run simultaneously. For the measurement of [$^{15}\text{N}_2$]urea enrichment, 2-hydroxypyrimidine derivative of urea was prepared as described by us previously (13). Standard urea solutions of known enrichment were run simultaneously.

The Ra of glucose and urea were calculated, assuming a single compartment model, during steady state by tracer dilution (14): $\text{Ra} = [(100/\text{Ep}) - 1] \times \text{I}$, where Ra is the rate of appearance in μmol or $\text{mg}/\text{kg} \cdot \text{min}^{-1}$, Ep is the enrichment of the tracer in the plasma, and I is the rate of infusion of the tracer in μmol or $\text{mg}/\text{kg} \cdot \text{min}^{-1}$. This equation assumes 100% enrichment of the infused tracers. The standard curves of known enrichment were also prepared assuming 100% enrichment.

All data are reported as mean \pm SD. Statistical analyses were performed on an IBM personal computer using SPSS software (SPSS Inc., Chicago, IL). Groups were compared using analysis of variance and pair-wise comparisons were carried out using Duncan's multiple range test.

RESULTS

All infants were born at term gestation, were stable, and did not have any perinatal problems (Table 1). As anticipated, the IDM were heavier and the SGA infants were lighter compared with AGA infants (analysis of variance, $p < 0.01$). All infants were studied during fasting. The infants designated "not fed" had not received any feeding since birth and were studied at an average of 3 h after birth. The previously fed infants had received their initial formula or breast feeding; however, full feeding had not yet been established. These infants also fasted, *i.e.* skipped one meal. Therefore, the urea and glucose kinetics were measured at approximately 8 h after the last meal. As shown by us previously, such a fast would correspond to approximately 16-18 h of fasting for an adult (15).

The plasma glucose concentrations were similar in the three groups (Table 2). No significant effect of initiation of feeding was observed in the normal infants or IDM. The urea N concentration was slightly lower in the SGA infants and higher in the IDM as compared with the normal infants. C-peptide concentrations were similar in normal infants and IDM, reflecting good control of maternal metabolism.

Glucose and urea turnover rates are shown in Tables 2 and 3. Isotopic steady state was observed in all infants between 2½ and 3 h. The Ra of glucose was lower in IDM as compared with normal infants soon after birth ($p = 0.01$); however, in the infants studied later, the glucose Ra was similar in the two groups. In the SGA infants, glucose Ra was lower (not significantly) than that in normal infants. The rate of urea synthesis in the term AGA infants was 5.6 mg N/kg \cdot h $^{-1}$ (range 2.4-8.9 mg N/kg \cdot h $^{-1}$ or 1.4-5.3 $\mu\text{mol}/\text{kg} \cdot \text{min}^{-1}$). Urea synthesis was slightly higher in the infants studied after the initiation of feeding. Of interest is the higher rate of urea synthesis in the IDM and the lower rate in the SGA infants. However, none of these measurements were significantly different from those in normal infants. Estimated protein oxidation from these data will correspond to 0.87 g/kg \cdot d $^{-1}$ for the term AGA infants, 1.1 g/kg \cdot d $^{-1}$ for the IDM, and 0.7 g/kg \cdot d $^{-1}$ for the SGA infants.

No correlation was observed between the rates of urea synthesis and the rate of glucose appearance, either in individual groups or in all the groups collectively.

DISCUSSION

The results of the present study show that the rate of urea synthesis or protein oxidation in the full-term infant in the period immediately after birth corresponds to 5.6 mg N/kg \cdot h $^{-1}$ or 0.8

Table 2. Glucose kinetics in newborn infant (mean \pm SD)

	Glucose (mmol/L)	C-peptide (μ g/L)	[² H ₂]glucose infusion rate (μ mol/kg·min ⁻¹)	Glucose Ra (μ mol/kg·min ⁻¹)
Normals				
Not fed* (11)	3.11 \pm 0.51	0.81 \pm 0.29	0.178 \pm 0.06	24.03 \pm 4.05
Previously fed* (8)	2.80 \pm 0.80	0.78 \pm 0.24 (n = 7)	0.224 \pm 0.077	23.34 \pm 6.45
IDM				
Not fed* (6)	2.75 \pm 0.43	0.98 \pm 0.38 (n = 5)	0.151 \pm 0.004	19.62 \pm 2.14†
Previously fed* (4)	3.62 \pm 1.37	1.12 \pm 0.40	0.150 \pm 0.013	28.88 \pm 9.90‡
SGA				
Previously fed* (4)	3.46 \pm 0.43	0.64, 0.58	0.147 \pm 0.007	17.71 \pm 3.34

* See Table 1 for explanation.

† $p = 0.01$ compared with fasting normal infants.

‡ $p = 0.05$ compared with fasting IDM (0.02 by WRS test).

Table 3. Urea kinetics in fasting newborn infant (mean \pm SD)

	BUN (mmol/L)	[¹⁵ N ₂]urea infusion rate (μ mol/kg·h ⁻¹)	Urea N Ra	
			mg N/kg·h ⁻¹	μ mol/kg·min ⁻¹
Normal				
Not fed* (11)	3.36 \pm 0.66	3.44 \pm 0.65	5.58 \pm 1.96	3.32 \pm 1.17
Previously fed* (8)	3.08 \pm 1.27	3.53 \pm 0.30	6.20 \pm 2.13	3.69 \pm 1.27
IDM				
Not fed* (6)	3.82 \pm 0.58	3.89 \pm 0.46	6.93 \pm 1.37	4.13 \pm 0.82
Previously fed* (4)	4.14 \pm 1.98	3.33 \pm 0.924	7.34 \pm 4.87	4.37 \pm 2.90
SGA				
Previously fed* (4)	2.37 \pm 0.86	3.79 \pm 2.11	4.60 \pm 1.23	2.74 \pm 0.73

* See Table 1 for explanation.

g/kg·d⁻¹ protein. Inasmuch as the babies had not been fed and were examined within 5 h of birth, these data represent the oxidation of endogenous protein. In addition, if one assumes that protein metabolism after birth is a continuum of fetal protein metabolism, these data may be the closest approximation of the rate of oxidation of protein by the human fetus at term gestation. These data are much lower than those estimated by Gresham *et al.* (3) based on the umbilical artery-vein gradient of urea in the human fetus at term gestation. The differences may be related to the fact that Gresham *et al.* only measured umbilical artery and vein differences at delivery and calculated urea production from the known clearance rate of urea by the sheep placenta. In the present study, the rate of urea synthesis was measured directly by tracer dilution. The ability of the human fetal liver to synthesize urea has been demonstrated as early as 12–16 wk of gestation (16), and significant activity of hepatic urea cycle enzymes has been measured at 13 wk of gestation (17). The fetus achieves 90% of the adult urea cycle enzyme activity by 36 wk of gestation. Thus, the human newborn at term has almost the full adult potential to synthesize urea.

The rate of protein oxidation is often quantified in newborn infants as well as in adults to estimate the N loss and, indirectly, N accretion and dietary protein requirements. The commonly used methods involve measurements of urinary N excretion (18, 19) or tracer isotope methods of quantifying oxidation to CO₂ of an essential amino acid (6, 7). The measurement of urinary N excretion rate is particularly difficult in infants and children because of the problems of accurate and complete collection of urine. In addition, it does not include nonurinary losses of N, *e.g.* skin and gut, does not include loss of urea as a result of hydrolysis in the gut (if any), and requires adjustments for the change in body urea pool size (4, 5, 20). The tracer carbon oxidation method also has the disadvantage of requiring the accurate measurement of the rate of carbon dioxide production, estimation of bicarbonate retention, and the effect of exchange of tracer carbon, resulting in a loss of tracer. In contrast, the

[¹⁵N₂]urea tracer dilution method includes all the urea synthesized by liver or appearing in plasma, irrespective of its source or the mechanism of loss. The use of doubly labeled tracer and measurement of the dilution of (m+2) species results in inclusion of all newly synthesized urea (10, 21, 22). Finally, the tracer (m+2) species do not include any recycled tracer, which will form only (m+1) species. As stated above, the tracer method includes all the urea appearing in plasma, irrespective of the source, and thus will also include dietary urea, particularly in the breast-fed infant because of the presence of a significant amount of urea N in breast milk (23–25). However, in the present study because all the infants were fasting, the contribution of urea N from diet was expected to be nil. Therefore, the present estimates of urea synthesis represent the best approximation of the synthesis of urea or protein oxidation of the newborn infants.

Only one other study has quantified rates of urea synthesis in newborn infants (26). Wheeler *et al.* (26), in four infants admitted to surgical intensive care units, by using orally administered [¹⁵N¹⁵N]urea and measuring ¹⁵N enrichment in urinary urea, estimated the urea synthesis rate to be 17.3 mmol N/kg·d⁻¹ or 12.0 μ mol of urea N/kg·min⁻¹, a rate considerably higher than that reported here. These differences may be related to the clinical condition of the infants, because all infants in our study were clinically well and had no perinatal problems. In addition, we studied infants during fasting, whereas the study population of Wheeler *et al.* was examined during feeding.

Other studies have quantified protein oxidation by measuring the rate of urea excretion or by the oxidation of essential amino acids. Pencharz *et al.* (27) estimated the obligatory loss of N in one full-term and six preterm infants from the regression of N balance and N intake. They estimated the total obligatory N loss to be 145 mg·kg⁻¹·d⁻¹ or 0.9 g protein kg⁻¹·d⁻¹, a rate similar to the measurements in the present study. Estimates of protein oxidation in full-term infants using [1-¹³C]leucine by us were ~1.4 g/kg·d⁻¹, which is much higher than that in the present study (4, 5, 19, 20). These differences may be related to the

observation that urea cycle enzymes can be readily regulated to conserve N. Furthermore, differences between protein oxidation measured by leucine tracer and that measured by urea excretion have also been observed under other physiologic circumstances. In only one study of adults, the rates of protein oxidation measured by [^{15}C]leucine tracer using keto-isocaproic acid enrichment and those measured by urinary urea N excretion were similar (28). Thus, the difference in other circumstances may be related to the regulation of urea synthesis or lack of measurement of the true precursor pool for the oxidation of amino acid.

The similar rate of urea synthesis in IDM as compared with normal infants is of interest, particularly in view of the observation of Jahoor and Wolfe (29) and others (30, 31) that glucose by itself or via its hormonal responses suppresses urea production. Our data may be related to the rigorous regulation of maternal metabolism so that these infants were no longer hyperinsulinemic. Finally, SGA infants synthesized less urea compared with the normal infants, although the differences were not statistically significant and the numbers of SGA infants studied were too few to draw a conclusion. The lower rate of urea synthesis in SGA infants may be related to the increased accretion of N as demonstrated in infants recovering from malnutrition (32). These data showed a decreased rate of urea excretion in infants recovering from malnutrition.

Because the major source of glucose Ra at this stage of fasting is hepatic glycogen and because gluconeogenesis from amino acid is only a small contributor, the lack of correlation between glucose Ra and urea synthesis rate was not surprising.

In summary, the present study quantified rates of urea synthesis during fasting in normal infants, IDM, and SGA infants. These data show that the rate of protein oxidation in newborn infants is $0.87 \text{ g/kg} \cdot \text{d}^{-1}$. The rate was slightly higher in IDM and lower in SGA infants. Because these observations were made during fasting either soon after birth, before any feeding, or after initiation of feeding, these data should be considered to reflect obligatory protein oxidation.

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