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Glycine, a Dispensable Amino Acid, Is Conditionally Indispensable in Late Stages of Human Pregnancy

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ABSTRACT

Background: Recently, we showed that there are higher protein, lysine, and phenylalanine requirements in late stages of pregnancy compared with early stages. Animal studies have suggested an increased dietary need for specific dispensable amino acids in pregnancy; whether such a need exists in human pregnancies is unknown.

Objective: The objective of the current study was to examine whether healthy pregnant women at midgestation (20–29 wk) and late gestation (30–40 wk) have a dietary demand for glycine, a dispensable amino acid, using the indicator amino acid oxidation method and measurement of plasma 5-oxoproline concentrations.

Methods: Seventeen healthy women (aged 26–36 y) randomly received different test glycine intakes (range: 5–100 mg·kg⁻¹·d⁻¹) during each study day in midgestation (~26 wk, n = 17 observations in 9 women) and late gestation (~35 wk, n = 19 observations in 8 women). Diets were isocaloric with energy at 1.7 × resting energy expenditure. Protein was given as a crystalline amino acid mixture based on egg protein composition at current estimated average requirement (EAR; 0.88 g·kg⁻¹·d⁻¹). Breath samples were collected at baseline and isotopic steady state to measure oxidation of L-[1–¹³C]phenylalanine to ¹³CO₂ (F¹³CO₂). Plasma was collected at the sixth hour of the study day. Linear regression crossover analysis and simple linear regression were used to assess responses in F¹³CO₂ and plasma 5-oxoproline concentrations to different glycine intakes.

Results: No statistically significant responses were observed in midgestation. However, in late gestation, lower glycine intakes resulted in higher rates of $F^{13}CO_2$ (suggesting low protein synthesis) with a breakpoint for phenylalanine oxidation at >37 mg glycine·kg⁻¹·d⁻¹ and higher plasma 5-oxoproline (suggesting low glycine availability) with a breakpoint >27 mg glycine·kg⁻¹·d⁻¹.

Conclusions: The findings suggest that glycine should be considered a "conditionally" indispensable amino acid during late gestation, especially when protein intakes are at 0.88 g·kg⁻¹·d⁻¹, the current EAR. This trial was registered at clinicaltrials.gov as NCT02149953. *J Nutr* 2021;151:361–369.

Keywords: glycine, pregnancy, requirements, human, stable isotopes

Introduction

Pregnancy is accompanied by changes in energy and nutrient requirements due to profound body adaptations and high rates of tissue synthesis (1). We determined the protein requirements during early (~16 wk) and late (~36 wk) gestation in healthy singleton pregnant women using the indicator amino acid oxidation (IAAO) method. The estimated average requirements (EARs) were determined to be 1.22 g·kg⁻¹·d⁻¹ (upper 95% CI: 1.66 g·kg⁻¹·d⁻¹) and 1.52 g·kg⁻¹·d⁻¹ (upper 95% CI: 1.77 g·kg⁻¹·d⁻¹) during early and late gestation, respectively (2). These values are considerably higher than the current EAR and RDA of 0.88 and 1.1 g·kg⁻¹·d⁻¹, respectively

(2, 3), although we have reported that current median intakes in healthy pregnant women are 1.3 and 1.5 $g \cdot kg^{-1} \cdot d^{-1}$ (3). Further, we recently demonstrated that the indispensable amino acid (IAA) lysine and phenylalanine requirements during late gestation are higher compared with early gestation (4, 5).

While the IAAs are clearly a major focus to ensure adequate dietary guidelines in pregnancy, the dispensable amino acids (DAAs) have recently been of focus in animal reproduction. Supplementing pig diets with arginine and/or glutamine has been shown to improve piglet birth weight and improve efficiency of nutrient utilization by optimizing placental growth (6). In a recent study, Tessari (7) estimated adult human DAA needs using factorial calculations. It was suggested that under

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some conditions, the endogenous synthesis of DAA may not be sufficient to meet body demands.

Glycine has traditionally been classified as a DAA, as it can be synthesized in the human body. Glycine is used for synthesis of glutathione, heme, creatine, nucleic acid, and uric acid (8). In addition, glycine is a main component of bile acids and makes up one-third of the amino acids in collagen, the most abundant protein in the human body (9). It has been suggested that de novo synthesis of glycine is insufficient to supply the metabolic demand (8, 9). Studies have shown decreased glycine flux in pregnant adolescents, suggesting an inability to maintain endogenous glycine production (10). With increasing stages of pregnancy, there is a progressive increase in excretion of 5-oxoproline (11). Conversion of γ -glutamylcysteine to glutathione (GSH) requires glycine, and when glycine availability is low, 5-oxoproline is formed (Supplemental Figure 1). Furthermore, glycine plays a central role in 1-carbon metabolism, and during pregnancy, perturbations of the transfer of methyl groups might affect cell proliferation and function (12).

Recently, we explored the role of 9 of the DAAs [alanine (Ala), arginine (Arg), asparagine (Asn), aspartate (Asp), glutamine (Gln), glutamate (Glu), glycine (Gly), proline (Pro), serine (Ser)] as an ideal nitrogen source to improve wholebody protein synthesis using the IAAO method in adults. We observed that 7 of the 9 DAAs (Ala, Arg, Asn, Asp, Glu, Gly, Ser) decreased IAAO significantly, except Gln and Pro (13). Whether such differences exist during pregnancy, especially during different stages of gestation, is unknown. We explored glycine's role in pregnancy first because of its implications in 1-carbon metabolism and because of the potential of using 5-oxoproline as a biomarker of glycine status. Therefore, the objectives of the current study were to determine whether there is a dietary demand for glycine during midgestation (20-29 wk) or late gestation (30–39 wk), examined using the IAAO, and measurement of plasma concentrations of 5-oxoproline and 1-carbon metabolites.

Methods

Participants

Healthy women aged between 20 and 40 y, pregnant with a single child, with reported prepregnancy BMI < 30 (in kg/m²) were recruited for the current study. All participants provided written and informed consent and were screened to ensure they were not experiencing severe nausea or vomiting, gestational diabetes, preeclampsia, or any other health conditions. The women were also interviewed about prescription medication and supplement intake. **Supplemental Figure 2** provides details on study flow, including screening and enrollment of pregnant participants. Financial compensation for transportation costs and an honorarium were offered to all participants upon completion of each study day. The study protocol was approved by the British Columbia Children's and Women's Hospital Research Ethics Board

Supplemental Table 1 and Supplemental Figures 1–4 are available from the "Online Supporting Material" link in the online posting of the article and from the same link in the online table of contents at http://jn.nutrition.org.

(H14-00495), and the study was registered at clinicaltrials.gov as NCT02149953.

Experimental design

The study design was based on the IAAO technique previously used to determine protein, lysine, and phenylalanine requirements in healthy pregnant women (2, 4, 5). Midgestation and late gestation were defined as 20–29 wk and 30–39 wk, respectively. The protocol included intakes ranging from 5 to 100 mg·kg⁻¹·d⁻¹, in 5-mg·kg⁻¹·d⁻¹ increments, selected randomly. Due to a combination of excluded study days (Supplemental Figure 2) and the selection method, not all intakes were used in each gestational stage. All study days completed by the same participant were separated by ≥ 5 d.

Preassessment

To determine eligibility, all potential participants attended a prestudy assessment at the Clinical Research and Evaluation Unit, BC Children's Hospital Research Institute. Participants arrived after a 10- to 12-h fast and were instructed to minimize physical exertion before the assessment. Fasted blood glucose concentrations were measured by finger prick (Ultra 2 LifeScan; One Touch) to screen for gestational diabetes, with a cutoff value of \geq 5.3 mmol/L, based on the guidelines from the Canadian Diabetes Association (14). Protein and glucose in urine were assessed using Chemstrip 7 Urinalysis Strips (Roche Diagnostics) as potential indicators of gestational diabetes and preeclampsia, respectively. Body composition was measured by 3-site (biceps, triceps, and subscapular) skinfold assessments (Harpenden Calipers; Baty International) with site, sex, gestation stage, and age-specific factors used (15, 16) to calculate fat-free mass. Resting energy expenditure (REE) was assessed by continuous, open-circuit indirect calorimetry (Vmax Encore). Each woman's height and weight were measured using a stadiometer and digital scale, respectively.

A brief medical and pregnancy history was collected to screen for medication use, pregnancy complications, and general health. Two days prior to each study day, participants were prescribed a standardization diet, recommending protein intakes at $1.5 \text{ g}\cdot\text{kg}^{-1}\cdot\text{d}^{-1}$. The prescribed diets were based on food sources preferred by each participant, as indicated by a 2-d food record collected during the prestudy assessment. Participants also kept a 2-d food record 2 d prior to coming in for the study, in order to assess protein intake.

Study day protocol

Each study day protocol is outlined in Figure 1. On the study day, participants arrived at the Clinical Research and Evaluation Unit after a 10- to 12-h fast. Height, weight, fasted blood glucose, and urinary measurements of glucose and protein were repeated at the beginning of each study day. Participants were randomly assigned to receive a test glycine intake (range: 5.0-100 mg·kg⁻¹·d⁻¹). The study day diet was consumed as 8-hourly isocaloric and isonitrogenous meals, each meal providing 1/12th of the daily energy requirement. The daily energy requirement was calculated as $1.7 \times \text{REE}$ for each participant, as well as protein at 0.88 $g kg^{-1} d^{-1}$, based on the current EAR for healthy pregnant women (17). Each meal consisted of a small protein shake and protein-free cookies. The shakes contained a liquid formula made with protein-free powder (PFD1; Mead Johnson Nutrition), flavored drink crystals (Tang and Kool-Aid; Kraft Canada, Inc.), and corn oil (Mazola; ACH Food Companies, Inc.). The macronutrient composition of the diet was \sim 53% from carbohydrate, \sim 38% from fat, and \sim 9% protein. The test protein was provided as a crystalline L-amino acid mixture (Ajinomoto, Inc.) based on the amino acid composition of egg protein, with the exception of glycine, and phenylalanine and tyrosine were provided at 31 and 61 mg kg⁻¹ d⁻¹, respectively. As previously described by Elango et al. (18), the presence of excess tyrosine is necessary to minimize retention of the ¹³C label in the tyrosine pool. This ensures partitioning of the carboxyl carbon from phenylalanine and thus incorporation into protein or oxidation (18). With the exception of water, participants did not consume anything besides the experimental diets during the study day.

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Abbreviations used: APE, atom percent excess; DAA, dispensable amino acid; DMG, dimethylglycine; EAR, estimated average requirement; F¹³CO₂, rate of appearance of (¹³C) labeled carbon dioxide in breath; GSH, glutathione; HFBA, heptafluorobutyric acid; IAA, indispensable amino acids; IAAO, indicator amino acid oxidation; REE, resting energy expenditure; SAH, *S*-adenosylhomocysteine; SAM, *S*-adenosylmethionine; SHMT, serine hydroxymethyltransferase.



FIGURE 1 Study day protocol. The experimental diets were given as hourly meals for 8 h. Priming doses of $NaH^{13}CO_3$ and L-[1-¹³C]phenylalanine were given with the fifth meal; hourly doses of L-[1-¹³C]phenylalanine were given with meals 5 through 8. Three breath samples were collected prior to the tracer protocol. Six plateau breath samples were collected after the tracer protocol. One venous blood sample was collected after the sixth hourly meal. Carbon dioxide production rate (VCO₂) was measured by indirect calorimetry after the fifth hourly meal. A fasting blood glucose measurement and a urine sample were taken before the first meal.

Tracer protocol

During each 8-h study day, the participants initially consumed 4-hourly meals not containing L- $[1-^{13}C]$ phenylalanine to allow for baseline sample collection. An oral priming dose of 0.264 mg·kg⁻¹ NaH¹³CO₃ (99 atom percent excess; Cambridge Isotope Laboratories, Inc.) and 4.0 mg·kg⁻¹ L- $[1-^{13}C]$ phenylalanine (99 atom percent excess; Cambridge Isotope Laboratories, Inc.) was provided at the fifth meal. Hourly oral doses of 3.0 mg·kg⁻¹·h⁻¹ L- $[1-^{13}C]$ phenylalanine were provided with the sixth to eighth meals until the end of the study (Figure 1).

Breath sample collection and analysis

During each study day, 3 and 6 breath samples were collected before (baseline) and after (isotopic steady state) the introduction of tracer amino acid, respectively, to measure the oxidation of L-[1–¹³C]phenylalanine to ¹³CO₂ ($F^{13}CO_2$). Breath samples were collected using disposable exetainer tubes (Labco Limited) and rate of carbon dioxide production was measured using open-circuit indirect calorimetry (V_{MAX} Encore; Viasys). Baseline breath samples were collected 45, 30, and 15 min before the tracer protocol began at hour 5, and isotopic steady-state samples were collected 150, 165, 180, 195, 210, and 240 min after the tracer protocol (Figure 1). Breath samples were stored at room temperature until analysis. Breath ¹³CO₂ enrichment was determined using continuous-flow isotope ratio mass spectrometry (Isoprime Ltd) and expressed as atom percent excess (APE) (19).

Plasma amino acids and metabolites

Glycine interacts with the methionine cycle and in metabolism with several 1-carbon metabolites (12) as outlined in Supplemental Figure 1. Therefore, plasma concentrations of related amino acids and metabolites were measured and compared between mid and late stages of pregnancy in response to glycine intakes.

Plasma collection

Venous blood samples were collected at BC Children's and Women's Hospital blood collection laboratory by a certified phlebotomist, using EDTA as the anticoagulant. The plasma was separated by centrifugation at 3000 \times g at 4°C for 10 min, sampled, and stored immediately at -80° C until analyzed. Plasma samples were collected at the sixth hour of the study to ensure a steady-state period of amino acid enrichment.

Plasma analytical methods

Plasma free amino acid concentrations were determined by ion exchange chromatography, using an amino acid analyzer (Hitachi L8900), as previously described (20). 5-Oxoproline was quantified by LC-MS/MS, as described previously (21). Choline, betaine, dimethylglycine (DMG), methionine, cysteine, and total homocysteine were analyzed by HPLC-MS/MS as previously described in detail by Friesen et al. (21) and Innis and Hasman (22).

S-adenosylmethionine (SAM) and S-adenosylhomocysteine (SAH) were analyzed by Waters H-class UHPLC and Waters Xevo tandem mass spectrometer (Waters Corp.), using a Zorbax SB-Aqua 2.1 × 100-mm column 3.5-micron particle size column with guard column (Agilent). Mobile phase A was composed of deionized water with 0.2% heptafluorobutyric acid (HFBA) and 0.1% formic acid. Mobile phase B consisted of methanol, LCMS grade; 0.2% HFBA; and 0.1% formic acid. Gradient separation was performed at a flow rate of 300 μ L/min starting at 100% A. Then, 50 μ L plasma was added to an Eppendorf tube containing 10 μ L internal standard. Next, 50 μ L 20% HFBA was added, and the sample was vortexed and left to stand at room temperature for 10 min, then centrifuged at 20,000 × g for 10 min at 4°C. The supernatant was removed and injected directly into the instrument.

Tracer oxidation

The rate of phenylalanine tracer oxidation ($F^{13}CO_2$, µmol·kg⁻¹·h⁻¹) was calculated as follows:

$$F^{13}CO_2 = (FCO_2)(ECO_2)(44.6)(60)/W(0.82)(100)$$
 (1)

where FCO₂ represents carbon dioxide production (mL/min), ECO₂ is the ¹³CO₂ enrichment in breath at plateau isotopic steady state (APE), W is the weight (kg) of the subject, 44.6 (μ mol/L) and 60 (min/h) are constants used to convert FCO₂ to μ mol/h, 0.82 is the correction factor for carbon dioxide retained by the body due to bicarbonate fixation, and 100 is used to convert APE to a fraction (23).

Statistical analysis

Results are reported as means \pm SDs. Breakpoint analysis was performed using a 2-phase linear regression crossover model of the F¹³CO₂, 5-oxoproline, SAM, and SAH data from late stages of pregnancy. The method selects the model with minimum residual standard error in a stepwise partitioning of glycine intake values (*x*) between 2 regression lines. The lines are estimated for each selected candidate breakpoint using mixed models (Proc Mixed, Statistical

TABLE 1	Characteristics of pregnant women a
preassessm	ent during mid and late gestation ¹

Observation in the	Midaestation	Late seated as
	windgestation	Late gestation
Participants, ² n	9	8
Gestational age, wk	23.3 ± 4.2	$31.8~\pm~2.3$
Age, y	$30.7~\pm~3.1$	$31.4~\pm~3.8$
Prepregnancy weight, kg	61.7 ± 13.2	$59.0~\pm~9.7$
Height, m	$1.64~\pm~0.08$	$1.62~\pm~0.06$
Prepregnancy BMI, ³ kg/m ²	$22.7~\pm~3.2$	$22.2~\pm~2.9$
Fat mass, ⁴ %	$26.7~\pm~4.9$	22.3 ± 3.1
Resting energy expenditure, ⁵ kcal/d	$1483~\pm~195$	$1484~\pm~210$

¹Values are means \pm SDs, n = 17 observations in 9 women (mid) or 19 observations in 8 women (late).

²Three women participated in both mid- and late-gestation studies.

³Based on participant reported prepregnancy weight.

⁴Determined by skinfold measurements (Harpenden Skinfold Caliper; Baty International)

⁵Determined by open-circuit indirect calorimetry (VMax Encore; Viasys).

Analysis Systems—SAS/STAT version 9.4; SAS Institute) to account for variability in number of completed study days per participant (24). The final model that best fit the data with the lowest SE, lowest root mean square error, and the highest R^2 identified the breakpoint. The 95% CI was calculated using Fieller's theorem (25): 95% CI = breakpoint $\pm t_{df, \alpha/2} \times$ SE, where SE is the SE of the combined regression lines, df is the degrees of freedom associated with the residual mean square of the best-fit model, and α is the 95% CI level (2, 19). The effect of glycine intake on plasma concentrations of the remaining biomarkers was assessed using linear regression (GraphPad Prism 6; GraphPad Software). Significance was set at P < 0.05 for all analysis.

Results

Participants

A total of 17 $(n_{mid} = 8 \text{ and } n_{late} = 9)$ women were studied, completing 36 individual study days (17 study days in midgestation and 19 study days in late gestation) (Table 1, Supplemental Figure 2). Three participants were studied in both midgestation and late gestation, each completing 2, 1, and 5 study days in midgestation and 2, 3, and 2 in late gestation, respectively (Supplemental Table 1). For these 3 women, a second prestudy assessment was performed prior to study days conducted during late gestation, to allow measurement for updated body weight and REE, as well as to reassess their health. All participants were otherwise unique between stages. Of the remaining participants recruited for midgestation, 3 participants came for 1 study day and 3 participants came for 2 study days. For late gestation among the remaining participants recruited, 1 participant came for 1 study day, 2 participants came for 2 study days, 1 participant came for 3 study days, and 1 participant came for 4 study days (Supplemental Table 1).

Women who participated in this study were not experiencing nausea or vomiting during the study days and reported no pregnancy complications. One woman reported having Crohn disease; however, the disease was inactive while participating in the study. Two women reported using SynthyroidTM (levothyroxine, AbbVie Inc.) for hypothyroidism and 1 woman reported using DiclectinTM (doxylamine succinate – pyridoxine, Duchesnay Inc.) for pregnancy-related nausea in midpregnancy. One woman had finished consuming antibiotics the day prior to coming in for a study day due to a urinary tract infection. No medications were consumed on the study day. None of the **TABLE 2** Study day assessments of healthy pregnant women during midgestation and late gestation¹

Variable	Midgestation $(n = 17)$	Late gestation (n = 19)
Weight, kg	64.9 ± 11.6	72.4 ± 10.4
Fasting blood glucose, mmol/L	4.4 ± 0.4	4.6 ± 0.5
VCO ₂ , mL/min	225 ± 32	257 ± 38
Gestational age, wk	25.9 ± 3.3	34.6 ± 2.7
Energy intake, kcal/d	2438 ± 338	2620 ± 335
Protein intake before study day, ² g·kg ⁻¹ ·d ⁻¹	1.2 ± 0.3	1.3 ± 0.3

¹Values are means \pm SDs, n = 17 observations in 9 women (mid) or 19 observations in 8 women (late). VCO₂, volume of carbon dioxide production per minute. ²Amount of protein consumed by participants in the 2 d before the study day, as indicated by dietary records.

women studied reported consuming alcohol or illicit substances at any time during their pregnancy. All participants were consuming daily prenatal multivitamin supplements for the duration of enrollment in this study.

The mean \pm SD prepregnancy BMIs were within normal range for both mid (22.7 \pm 3.2) and late (22.2 \pm 2.9) pregnancy stages (26), and fasting blood glucose concentrations were \leq 5.3 mmol/L (Table 2) (14, 26). The mean \pm SD protein intakes for the 2 d prior to each study day were lower than prescribed (1.2 \pm 0.3 and 1.3 \pm 0.3 g·kg⁻¹·d⁻¹ for midgestation and late gestation, respectively) but similar to our recent pregnancy study (5).

L-[1–¹³C]phenylalanine oxidation

With increasing intakes of glycine $(5-100 \text{ mg}\cdot\text{kg}^{-1}\cdot\text{d}^{-1})$ in midgestation, linear regression analysis showed no significant relation ($R^2 = 0.127$, P = 0.161; Figure 2A). In late gestation, the oxidation of L-[1–¹³C]phenylalanine (F¹³CO₂) was higher with low glycine intakes and declined with increasing glycine intakes. Two-phased linear regression analysis of the F¹³CO₂ data identified a breakpoint at 37 mg·kg⁻¹·d⁻¹($R^2 = 0.57$; 95% CI: 17, 58 mg·kg⁻¹·d⁻¹; Figure 2C).

5-Oxoproline

With increasing glycine intakes $(5-100 \text{ mg}\cdot\text{kg}^{-1}\cdot\text{d}^{-1})$ in midgestation, no pattern was observed in plasma 5-oxoproline concentrations ($R^2 = 0.065$, P = 0.322; Figure 2B). However, plasma 5-oxoproline concentrations decreased with increasing intakes of glycine ($5-100 \text{ mg}\cdot\text{kg}^{-1}\cdot\text{d}^{-1}$) in late gestation (Figure 2D). Two-phased linear regression analysis identified a breakpoint at 27 mg $\cdot\text{kg}^{-1}\cdot\text{d}^{-1}$ ($R^2 = 0.31$; 95% CI: 14, 40 mg $\cdot\text{kg}^{-1}\cdot\text{d}^{-1}$; Figure 2D).

Plasma concentrations of amino acids and metabolites

Glycine and serine.

During midgestation, plasma glycine and serine concentrations increased significantly with increasing glycine intakes (5–100 mg·kg⁻¹·d⁻¹) ($R^2 = 0.667$, P = 0.0001 and $R^2 = 0.437$, P = 0.0053, respectively; Figure 3A,B), whereas no significant increase in plasma glycine or serine concentrations was observed in late gestation ($R^2 = 0.117$, P = 0.152 and $R^2 = 0.045$, P = 0.381, respectively; Figure 3C,D).

Methionine, homocysteine, and cysteine.

No significant change in plasma methionine concentration was observed for either midgestation ($R^2 = 0.007$, P = 0.759;



FIGURE 2 Effect of graded glycine intakes on F¹³CO₂ (A, C) and plasma 5-oxoproline (B, D) during midgestation (A, B) and late (C, D) gestation in healthy pregnant women. There were 17 observations in 9 women (mid) or 19 observations in 8 women (late). Linear regression analysis was performed on midgestation data. Biphase linear regression crossover analysis was performed on late gestation data.

Figure 4A) or late gestation ($R^2 = 0.017$, P = 0.598; Figure 4D) with increasing glycine intakes (5–100 mg·kg⁻¹·d⁻¹). Similarly, no significant changes in plasma concentrations of cysteine were observed for either midgestation or late gestation ($R^2 = 0.009$, P = 0.716 and $R^2 = 0.179$, P = 0.071, respectively; Figure 4C,F). Interestingly, plasma concentrations of homocysteine decreased significantly with increasing intakes of glycine (5–100 mg·kg⁻¹·d⁻¹) in late gestation ($R^2 = 0.197$; Figure 4E), which was not observed in midgestation ($R^2 = 0.148$, P = 0.127; Figure 4B).

SAM and SAH.

SAM and SAH concentrations decreased with increasing intakes of glycine (5–100 mg·kg⁻¹·d⁻¹) in late gestation. Twophased linear regression analysis identified breakpoints at 36 mg·kg⁻¹·d⁻¹ (95% CI: 21, 51 mg·kg⁻¹·d⁻¹; $R^2 = 0.37$; Figure 5C) for SAM and at 28 mg·kg⁻¹·d⁻¹ (95% CI: 22, 35 mg·kg⁻¹·d⁻¹; $R^2 = 0.60$; Figure 5D) for SAH in late gestation. In contrast, for midgestation, there were no significant changes in plasma concentrations of SAM ($R^2 = 0.002$, P = 0.857) or SAH ($R^2 = 0.15$, P = 0.13), with increasing glycine intakes (5–100 mg·kg⁻¹·d⁻¹) (Figure 5A,B).

Choline, betaine, DMG, and sarcosine.

There were no significant responses in plasma choline, betaine, or sarcosine to increasing glycine intakes $(5-100 \text{ mg} \cdot \text{kg}^{-1} \cdot \text{d}^{-1})$ during midgestation ($R^2 = 0.0005$, P = 0.934, $R^2 = 0.0003$;

P = 0.957, $R^2 = 0.046$, P = 0.425; Supplemental Figure 3A,B,D) and late gestation ($R^2 = 0.137$, P = 0.119, $R^2 = 0.005$; P = 0.772, $R^2 = 0.08$, P = 0.239; Supplemental Figure 3E,F,H). In midgestation, DMG concentrations tended to increase ($R^2 = 0.22$, P = 0.055), which was not observed in late gestation ($R^2 = 0.048$, P = 0.36) (Supplemental Figure 3C,G).

Ornithine, citrulline, arginine, urea, and histidine.

In both midgestation and late gestation, no significant changes were observed in plasma citrulline ($R^2 = 0.196$, P = 0.086 and $R^2 = 0.059$, P = 0.318; **Supplemental Figure 5B**,G) and arginine ($R^2 = 3.14\text{E-}07$, P = 0.998 and $R^2 = 0.012$, P = 0.657; Supplemental Figure 5C,H) in response to increased glycine intakes (5–100 mg·kg⁻¹·d⁻¹) in both midgestation and late gestation. However, ornithine ($R^2 = 0.289$, P = 0.018), histidine ($R^2 = 0.258$, P = 0.027), and urea ($R^2 = 0.543$, P = 0.003) significantly decreased with increasing glycine intakes in late gestation (**Supplemental Figure 4F**,J,I) but not in midgestation ($R^2 = 0.002$, P = 0.888, $R^2 = 0.0002$; P = 0.962, $R^2 = 0.004$, P = 0.807; Supplemental Figure 4A,E,D).

Discussion

To our knowledge, this is the first study to directly address the impact of a range of glycine intakes (low to high) in pregnant women at 2 distinct gestational stages (mid and late). Our



FIGURE 3 Plasma concentrations of glycine (A, C) and serine (B, D) in response to graded intakes of glycine during midgestation (A, B) and late gestation (C, D) in healthy pregnant women. There were 17 observations in 9 women (mid) or 19 observations in 8 women (late). Linear regression analysis was performed on all data.

results suggest that restriction of glycine in acute dietary studies during midgestation (~26 wk) does not have significant effects on whole-body protein synthesis, plasma 5-oxoproline, and 1-carbon metabolite concentrations. However, in late stages of gestation (~35 wk), restriction of glycine reduced wholebody protein synthesis (as observed by high ¹³C-phenylalanine oxidation, F¹³CO₂) and increased plasma concentrations of 5-oxoproline, which plateaued with increasing intakes of glycine (>37 mg·kg⁻¹·d⁻¹). Earlier 5-oxoproline excretion has been suggested as a marker for glycine status in pregnant women (27). Synthesis of GSH, a tripeptide of cysteine, glutamate, and glycine, is initiated when glutamate is combined with cysteine to form γ -glutamyl cysteine (Supplemental Figure 1). With the addition of glycine to γ -glutamyl cysteine, GSH is produced. However, when glycine is limiting, γ -glutamyl cysteine is metabolized to 5-oxoproline. Previous studies have shown that in late pregnancy, plasma 5-oxoproline is increased by 128% compared with nonpregnant women, possibly due to glycine insufficiency, which may limit GSH synthesis (21). Furthermore, Jackson et al. (11) showed that compared with nonpregnant women, urinary excretion of 5-oxoproline in late pregnancy was 365% higher when standardized against creatinine, suggesting that endogenous synthesis of glycine may be insufficient during late pregnancy. To our knowledge, our findings are the first to provide evidence of a direct relation between glycine intake and plasma 5-oxoproline concentrations during pregnancy. Taken together, our results suggest that glycine is conditionally indispensable during late stages of pregnancy.

intakes are associated with increased plasma SAM and SAH concentrations, which plateaued with increasing glycine intakes (>37 mg·kg⁻¹·d⁻¹). The exact mechanisms behind these findings are unknown. We hypothesize that with our study design supplying adequate methionine (and most 1-carbon nutrients involved in the methionine cycle) with low glycine intakes, it is possible that glycine N-methyltransferase, which catalyzes glycine to sarcosine, is downregulated (28). However, phosphatidylethanolamine N-methyltransferase, the enzyme responsible for converting phosphatidylethanolamine to phosphatidylcholine, which also converts SAM to SAH, is upregulated (29, 30). Similar to our results, recently, van Riet et al. (31) followed a set of sows throughout gestation and observed that by late stages of gestation, plasma concentrations of SAM and SAH were positively correlated with plasma methionine concentrations and negatively correlated with plasma glycine concentrations. Finkelstein (32) has stated earlier that plasma concentrations of SAM and SAH need to be interpreted with caution, as they do not reflect intracellular concentrations. However, since we observed changes in SAM and SAH plasma concentrations in late gestation and not in midgestation in response to glycine intakes using the same study design, the relative differences are comparable and suggest that glycine availability is insufficient in later stages of pregnancy.

In late but not midpregnancy, we found that low glycine

Structurally, glycine is the simplest amino acid and is a key component of proteins, such as collagen, and important for nucleic acids, heme, and creatine synthesis (8). Thus, a developing fetus has an increased demand for glycine.



FIGURE 4 Plasma concentrations of methionine (A, D), homocysteine (B, E), and cysteine (C, F) in response to graded intakes of glycine during midgestation (A, B, C) and late gestation (D, E, F) in healthy pregnant women. There were 17 observations in 9 women (mid) or 19 observations in 8 women (late). Linear regression analysis was performed on all data.

Previously, how the fetus meets the demand for glycine has been explored (12). Glycine can be synthesized from serine via serine hydroxymethyltransferase (SHMT), and since sheep placenta express a significant amount of SHMT, it was hypothesized that maternal serine is converted to glycine in the placenta and transferred to the fetus (33). But this theory has been questioned due to low SHMT activity in human placentas, and studies of amino acid uptake and interconversions suggest that the serineglycine transfers may not be that significant (34). In the current study, plasma concentrations of glycine and serine increased significantly with increasing glycine intakes in midgestation but not in late gestation. The women received adequate serine in the diet, and thus our results suggest that in human late-stage pregnancies, de novo synthesis of glycine was inadequate and there might be a demand for maternal preformed glycine.

From a physiologic viewpoint, all of the amino acids that occur in proteins, whether they are synthesized in the body or not, are essential for tissue protein synthesis and various other functions (35). Recently, there have been several discussions on whether a dietary need can be determined for DAA (7, 36, 37). Tessari (7) uses the term nonessential amino acid usage, and based on factorial estimates of wholebody amino acid composition and obligatory nitrogen losses, has suggested estimates for all amino acids. Glycine has been suggested to have a usage of 46 to 59 mg·kg⁻¹·d⁻¹ in healthy nonpregnant adults (7). Our results suggest that during late stages of pregnancy, there is a potential need for preformed glycine at 37 mg·kg⁻¹·d⁻¹. One key aspect in the current study is that the protein intake on study days was at the current recommended EAR of 0.88 g·kg⁻¹·d⁻¹ for protein during pregnancy (17). This is considerably lower than the EAR during late stages of pregnancy, as determined recently by us $(1.52 \text{ g}\cdot\text{kg}^{-1}\cdot\text{d}^{-1})$ (2). The women in our study received protein at 57-67 g/d and 64-73 g/d in midgestation and late gestation, respectively. These are lower than normally reported by pregnant mothers in Canada (~ 99 g/d) (3). Consequently,

while the moderate protein intake in the current study during midgestation did not affect de novo glycine synthesis, in late gestation, glycine synthesis was not sufficient. On average, highquality protein diets provide \sim 33 mg/kg protein, suggesting that an intake of ~ 1.1 g·kg⁻¹·d⁻¹ would be needed to satisfy a glycine need of 37 mg·kg⁻¹·d⁻¹ in pregnancy. While these intakes are normally achieved in developed countries (3), our findings have implications for pregnant populations who consume diets with low protein quantity and quality (38, 39). In developing countries, total protein intakes in pregnancy is ~50 g/d, which translates to ~0.83 g·kg⁻¹·d⁻¹ for a 60-kg body weight (38). Thus, glycine needs are unlikely to be met at these protein intakes, emphasizing the "conditional essential" nature of this amino acid. Similar to our study, Yu et al. (40), using ¹⁵N-tracers, showed earlier in adults that when dietary nitrogen was provided as IAA alone with a total protein at 0.6 $g \cdot kg^{-1} \cdot d^{-1}$ (EAR for adults) (17), alanine synthesis was not affected, but glycine synthesis declined (40). A later longer-term study in young adults, using L-5-[1-¹³C]oxoproline and diets either free of glycine or methionine + cysteine, showed by day 6 significantly higher urinary excretion and oxidation of 5-oxoproline (41). Thus, even in well-nourished adult men, glycine supply is crucial in the context of length of dietary restriction or when restricted in dispensable amino acid nitrogen.

Our study has a few limitations with respect to small sample size and the acute study design. However, the strength of the study is the use of a study design earlier applied during pregnancy to determine protein/amino acid requirements and measurement of several plasma metabolites related to glycine metabolism at 2 gestational stages in pregnancy (2, 4, 5). It should be emphasized, however, that our objective in the study was not to determine a "requirement" for glycine during pregnancy but rather test the hypothesis whether a dietary need for glycine exists during stages of pregnancy. Future studies need to be conducted with a range of glycine intakes with adequate



FIGURE 5 Plasma concentrations of *S*-adenosylmethionine (SAM) (A, C) and *S*-adenosylhomocysteine (SAH) (B, D) in response to graded intakes of glycine in healthy pregnant women during midgestation (A, B) and late gestation (C, D) in healthy pregnant women. There were 17 observations in 9 women (mid) or 19 observations in 8 women (late). Linear regression analysis was performed on midgestation data. Biphase linear regression crossover analysis was performed on late gestation data.

protein and measurement of other markers, including synthesis of glutathione and creatine during pregnancy (42).

In summary, the current study showed an increased rate of IAAO when glycine intakes were low, suggesting low protein synthesis in late gestation but not in midgestation. Plasma 5-oxoproline concentrations, an independent biomarker of glycine status, were also elevated with low glycine intakes only in late stages of pregnancy. Patterns of response in plasma 1-carbon metabolites, including SAM, SAH, and, to a lesser extent, choline, were also different in late gestation. Taken together, these findings suggest that glycine de novo synthesis may be insufficient during late pregnancy and is conditionally indispensable, especially when protein intakes are at the current recommended pregnancy EAR of $0.889 \cdot kg^{-1} \cdot d^{-1}$.

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