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Evaluation of Protein Quality in Humans and Insights on Stable Isotope Approaches to Measure Digestibility – A Review

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ABSTRACT

The recent Food and Agricultural Organization/World Health Organization/United Nations University expert consultations on protein requirements and quality have emphasized the need for the new Digestible Indispensable Amino Acid Score (DIAAS), as a measure of protein quality. This requires human measurements of the true ileal digestibility of individual indispensable amino acids (IAAs) until the end of the small intestine. Digestibility is measured using standard oro-ileal balance methods, which can only be achieved by an invasive naso-ileal intubation in healthy participants or fistulation at the terminal ileum. Significant efforts have been made over the last 2 decades to develop noninvasive or minimally invasive methods to measure IAA digestibility in humans. The application of intrinsically labeled (with stable isotopes like ¹³C, ¹⁵N, and ²H) dietary proteins has helped in circumventing the invasive oro-ileal balance techniques and allowed the differentiation between endogenous and exogenous protein. The noninvasive indicator amino acid oxidation (IAAO) technique, which is routinely employed to measure IAA requirements, has been modified to estimate metabolic availability (a sum of digestibility and utilization) of IAA in foods, but provides an estimate for a single IAA at a time and is burdensome for participants. The recently developed minimally invasive dual isotope tracer method measures small intestinal digestibility of multiple amino acids at once and is suitable for use in vulnerable groups and disease conditions. However, it remains to be validated against standard oro-ileal balance techniques. This review critically evaluates and compares the currently available stable isotope-based protein quality evaluation methods with a focus on the digestibility and metabolic availability measurements in humans. In view of building a reliable DIAAS database of various protein sources and subsequently supporting protein content claims in food labeling, a re-evaluation and harmonization of the available methods are necessary. *Adv Nutr* 2021;00:1–13.

Statement of Significance: This review is the first of its kind that exhaustively reports on, and critically compares, stable isotope-based and direct measurements of amino acid digestibility/bioavailability in humans, using the ileal balance, dual isotope, and IAAO methods. The review provides details on the principles, advantages, and drawbacks of different methods, as well as details on efficient approaches of intrinsic labeling of food proteins; in addition, this review details all available human measurements of IAA digestibility and metabolic availability of food proteins.

Keywords: protein quality, stable isotopes, protein digestibility, metabolic availability, intrinsic labeling, oro-ileal balance, dual isotope tracer technique, indicator amino acid oxidation

Introduction

Dietary protein provides nitrogen and amino acids (AAs), particularly indispensable amino acids (IAAs), which are required in adequate quantity and proportion for the synthesis of protein and other nitrogen- and AA-related compounds with various structural and biological functions in the body

(1, 2). The quality of dietary protein source has been directly assessed by measuring the utilization and retention of dietary nitrogen and AAs in the body, but this approach is difficult due to the complexity of the physiological and metabolic processes of protein digestion, absorption, and metabolic utilization of AAs (2–5). Alternatively, the quality

of a dietary protein is defined by its ability to meet age-specific nitrogen and IAA requirements for growth and maintenance. Therefore, it can be assessed by the widely accepted chemical scoring approach that compares the IAA pattern of a protein with the age-specific IAA requirements corrected for protein or IAA digestibility, through 2 simple indexes, the Protein Digestibility Corrected Amino Acid Score (PD-CAAS) and the Digestible Indispensable Amino Acid Score (DIAAS) (2, 5–7). A critical aspect of these indexes is measurement of protein and IAA digestibility to correct the chemical score. Several methods for measuring digestibility are currently available, where stable isotope labeling, especially intrinsically labeled dietary proteins with ^{15}N , ^2H , and ^{13}C tracers, has been used. This review critically evaluates and compares the methodological concerns of the stable isotope-based protein quality evaluation methods with a focus on the digestibility and metabolic availability (MA) measurements in humans and their applicability in various pathophysiological conditions.

Concepts of chemical scoring and correction for digestibility

The quality of a dietary protein can be assessed by its chemical score, which is a ratio of its IAA content to the age-specific IAA requirement pattern (4, 6, 7). The lowest chemical score of a food is corrected for the crude protein digestibility to obtain a protein quality metric called the PD-CAAS (**Supplementary Material 1**). Although practical and widely used, the PD-CAAS has been subject to criticism, mainly for using a single fecal nitrogen digestibility value for correction. Indeed, fecal digestibility is not always a good proxy of digestibility, especially for proteins of low digestibility (8), possibly due to the contribution of colonic microbes to nitrogen transactions via the fermentation of undigested protein entering the colon. As the PD-CAAS is truncated to 100% this does not allow indication of the potential of a high-quality protein to optimize the AA composition of food mixtures with low protein quality. The advantages and limitations of the PD-CAAS have been extensively reviewed previously (9–11). To overcome the concerns related to PD-CAAS, the World Health Organization/Food and Agriculture Organization/United Nations University (WHO/FAO/UNU) expert consultations recommended that the chemical score of each IAA should be corrected for their true ileal digestibility values measured in humans, and the lowest score thus obtained be termed the DIAAS (**Supplementary Material 1**) (4, 6). Unlike the PD-CAAS, a score of >100% for a single

food or mixed diets in the DIAAS is not truncated to indicate the potential of a high-quality protein to complement low-quality protein in mixed diets. Both the DIAAS and PD-CAAS can be used to inform protein content claims in food labeling. Since measuring true ileal AA digestibility in humans is invasive and expensive, pigs and alternatively rats are used to determine the ileal digestibility coefficients for regulatory purposes (4, 6, 12). The reliability of the PD-CAAS and DIAAS is also dependent on factors such as the determination of AA composition, the reference AA profile, the nitrogen to protein conversion factor, and the complexities and uncertainties around the measurements of dietary protein/IAA digestibility.

Digestibility issues: fecal versus ileal digestibility, apparent versus true digestibility, protein versus AA digestibility

The simplest and long-standing method of measuring protein and AA digestibility has been the oro-fecal balance method (5, 7). Although noninvasive, a major concern of the fecal digestibility measurements is the hindgut microbial modification of the undigested dietary nitrogen exiting the terminal ileum. As dietary nitrogen and AA absorption essentially occurs in the small intestine, the ileal digestibility, measured at the terminal ileum is considered to be a more accurate assay. The ileo-fecal differences in nitrogen (2–9%) and AA digestibility (0.4–15%) have been reported in monogastric animals (including humans) for highly digestible proteins (13, 14), and these differences were reported to be as high as 20% in rats for less digestible plant proteins possibly due to microbial fermentation of dietary fiber (15) and undigested AA during colonic transit (16).

The traditional assessment of digestibility is based on the measurements of total nitrogen and AA losses (endogenous and exogenous) in digesta and is termed “apparent” digestibility (9, 17, 18). Different methods have been used to measure endogenous losses; the advantages and limitations of these methods have been previously discussed (17, 19). When apparent digestibility is corrected for endogenous protein and AA losses measured by feeding with protein-free diets (20, 21) or by differentiating between endogenous and exogenous losses using intrinsically labeled proteins (22), it is termed “true” and “real” digestibility, respectively (**Supplementary Material 1**) (19). For simplicity, the term “true” is often used in place of “real” digestibility.

An additional concern in the assessment of dietary protein digestibility is the uncertainty associated with assuming overall protein (nitrogen) digestibility as a proxy for individual AA digestibility. A modest variation in ileal digestibility of IAAs has been reported in humans (14, 20, 22, 23) ranging from 89% (threonine) to 95% (lysine) with a nitrogen digestibility of 94% in soy protein isolate (23). Considerable differences have been observed in less digestible whole-plant protein sources, such as pea cultivars, where ileal digestibility of IAAs varied from 75% (tryptophan) to 89% (methionine) with a nitrogen digestibility of 76% in pigs (24), suggesting

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Supplemental Material 1 and Supplemental Table 1 are available from the “Supplementary data” link in the online posting of the article and from the same link in the online table of contents at <https://academic.oup.com/advances/>.

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Abbreviations used: AA, amino acid; DIAAS, Digestible Indispensable Amino Acid Score; IAA, indispensable amino acid; IAAO, indicator amino acid oxidation; MA, metabolic availability; NPPU, net postprandial protein utilization; PD-CAAS, Protein Digestibility Corrected Amino Acid Score; PPU, postprandial protein utilization; TAAD, true amino acid digestibility.

TABLE 1 Approaches used for intrinsic labeling of proteins to measure protein and AA digestibility in humans

Proteins	Label	Form of label	Method of administration
Milk and meat from ruminants	¹⁵ N	¹⁵ N ammonium sulphate	<ul style="list-style-type: none"> • Perfusion in rumen (22, 23, 26, 28) • Oral administration of dose (23, 78)
	² H	² H-labeled crystalline AAs Deuterium oxide (² H ₂ O)	<ul style="list-style-type: none"> • Intravenous infusion in jugular vein (98) • Oral administration of dose dissolved in water (38)
	¹³ C	¹³ C labeled crystalline AAs	<ul style="list-style-type: none"> • Feeding with ²H-labeled fodder (43) • Intravenous infusion in jugular vein (87, 91, 99)
Egg and chicken meat	¹⁵ N, ² H, and ¹³ C	¹⁵ N-, ² H-, and ¹³ C- labeled crystalline AAs	<ul style="list-style-type: none"> • Oral administration of dose dissolved in water (59) • Diets supplemented with labeled AAs (100–103)
Plant proteins	¹⁵ N	¹⁵ N ammonium nitrate, ¹⁵ N potassium nitrate, ¹⁵ N ammonium chloride	<ul style="list-style-type: none"> • Fertilization of soil with labeled salt (29, 35, 104) • Foliar spraying before flowering (23, 30, 80, 105)
	² H	Deuterium oxide (² H ₂ O)	<ul style="list-style-type: none"> • Pulse dosing at flowering (37, 59)
	¹³ C	¹³ CO ₂	<ul style="list-style-type: none"> • Atmospheric labeling (32, 45)

AA, amino acid.

the need to measure the digestibility of each IAA to evaluate the quality of dietary protein.

Intrinsic labeling of dietary protein for measuring protein and AA digestibility in humans

Intrinsically labeled dietary proteins with stable isotopes have been used to measure true or real digestibility for the last 25 y. Intravenous, oral, or ruminal administration of single/multiple labeled free AAs or ¹⁵N ammonium sulphate have been used for labeling milk (25, 26), eggs (27), and meat (28), whereas deuterium oxide (²H₂O) and ¹⁵N fertilizers (29–31), and more rarely ¹³CO₂ (32), have been used to intrinsically label plant proteins (Table 1). For milk proteins, the intravenous administration of labeled AAs is more efficient compared with oral administration as it avoids losses due to fermentation in the rumen, feed refusals, and impaired absorption due to the feed matrix. However, due to the high cost, the protein is generally labeled with 1 (33) or 2 AAs (34). Another consideration relates to the type of tracer being used, which depends on the method (direct or indirect) of determining digestibility. For instance, ¹⁵N-labeled milk, bovine meat, and plant proteins (lupin, soybean, wheat, rapeseed, and pea) have been widely used to measure ileal nitrogen and AA digestibility using direct ileal balance methods in humans (23, 28, 30, 31, 35, 36). However, due to the exchange and loss of ¹⁵N during transamination of AAs (Supplementary Material 1), the use of this label has major limits especially for the indirect methods which measure plasma appearances of labeled AAs (25, 37).

Goat milk protein has been previously labeled with ²H by administering ²H₂O to a lactating goat (38). In this method, labeling occurs by the exchange between the hydrogen atoms at the α-H position of AAs with ²H from enriched body water during the transamination reaction, with a small quantity of ²H labeled AAs being synthesized de novo (39–42). Similar to ¹⁵N, AAs can lose the α-²H label during the same reaction,

underestimating the absorbed ²H AAs (39). ²H-labeled milk and meat could also be produced by feeding with ²H-labeled fodders, obtained by watering fodder plants with ²H₂O (43). Since plants are autotrophs and synthesize AAs de novo, this would lead to ²H labeling in multiple random positions of the AAs, including α-H (44). Although use of ²H₂O enables efficient labeling of plant proteins, labeling milk using ²H-labeled fodder is cumbersome, expensive, and leads to lower enrichments in milk particularly for the limiting AAs in the plant.

The transamination losses of the ²H label can be addressed by using correction factors determined by controlled experiments (37) (Table 2, Supplementary Material 1). Transamination correction factors (F_{TCF}) have also been proposed for ¹⁵N but experimental validation is necessary due to their high intraindividual variability and the estimates differed with the amount of protein in test meals (25). This could be attributed to the higher potential for exchange of ¹⁵N, owing to a smaller pool of amino nitrogen compared with hydrogen in the body water (25, 37). Therefore, the best isotopic label to use is ¹³C, as this is not modified by transamination or other metabolic processes, and intrinsic labeling could be achieved by oral administration of ¹³C-labeled AAs to animals, or by growing plants/algae in a

TABLE 2 Transamination correction factors for IAAs using ²H¹

IAA	² H label
Methionine	1.058 ± 0.005
Phenylalanine	1.053 ± 0.006
Threonine	1.016 ± 0.002
Lysine	1.002 ± 0.002
Leucine	1.081 ± 0.002
Isoleucine	1.070 ± 0.004
Valine	1.048 ± 0.003

IAA, indispensable amino acid.

¹Values are mean ± SD, n = 6; Reproduced with permission from Devi et al. (37).

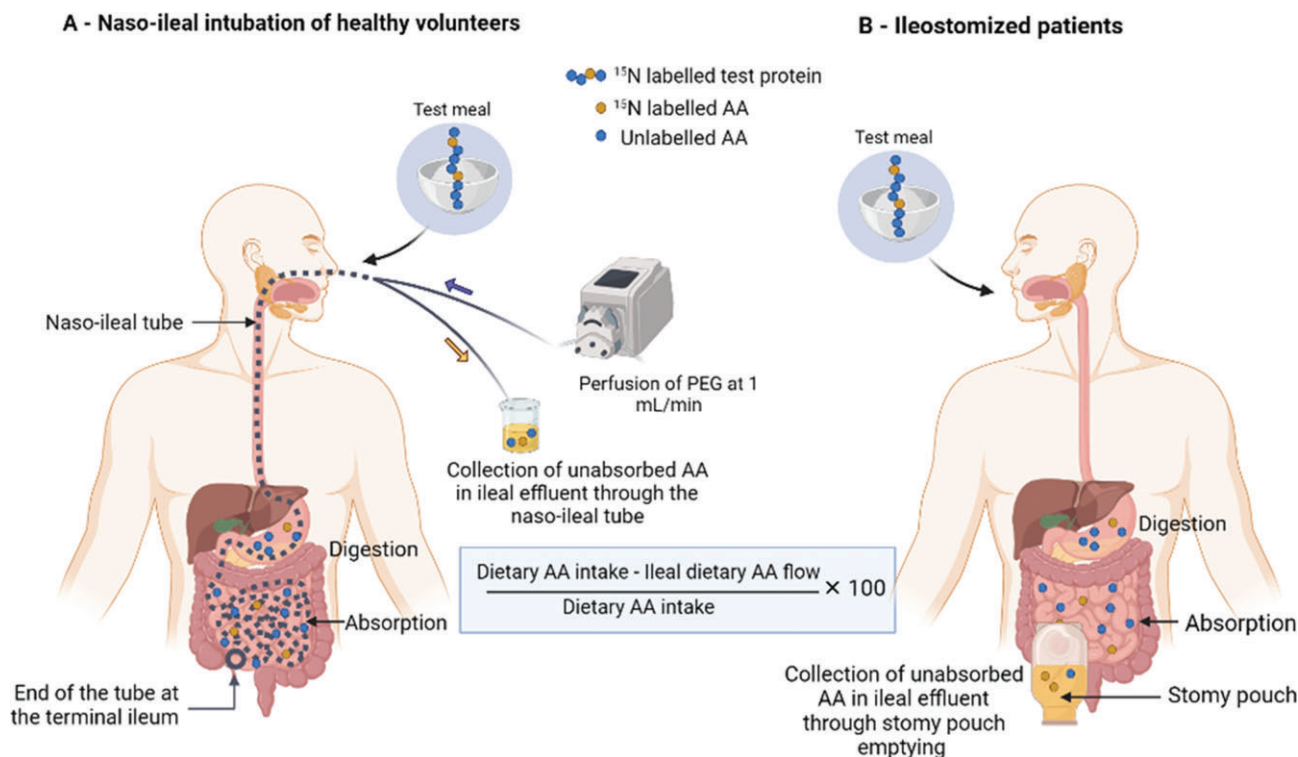


FIGURE 1 Schematic representation of the principle of the oro-ileal balance method to measure ileal AA digestibility with (A) intubated healthy volunteers or with (B) ileostomized patients using labeled protein. With the intubation method, the volunteers are equipped with a triple lumen naso-ileal tube. One lumen is dedicated to inflate a balloon at the end of the tube to facilitate the migration of the tube through the intestinal tract. When the tube is at the terminal ileum, a perfusion of PEG (unabsorbable marker) is started through the second lumen, in order to evaluate intestinal flow by the slow marker method. The test meal containing intrinsically ^{15}N -labeled test protein undergoes digestion and absorption and ileal effluents containing nonabsorbed dietary AAs are continuously collected from the third lumen during the 8-h postprandial period. In ileostomates, where the colon and rectum have been partially or totally removed and the terminal ileum exteriorized, the ileal effluents are directly collected into the pouch. The digestibility of each AA is determined by the ratio of the absorbed AA to the intake. AA, amino acid; PEG, polyethylene glycol.

$^{13}\text{CO}_2$ -enriched environment (32, 45). Although ^{13}C labeling is feasible for animal proteins, this might not be the case for plant proteins, as it is difficult and expensive to obtain the high grain yields required for digestibility protocols (46). Overall, it is important to choose an appropriate tracer for intrinsic labeling of dietary proteins based on the method used for measuring digestibility with careful consideration of the labeled atom position to avoid possible label losses due to metabolic rearrangements.

Direct method for true ileal protein and AA digestibility: oro-ileal balance method

The classical and standard method for measuring true ileal protein and AA digestibility entails collection of the digesta at the terminal ileum during the postprandial period to determine the amount of undigested nitrogen or/and AA. In humans, this requires invasive procedures unless studied specifically in ileostomates (20). In healthy volunteers with an intact intestine, a radio-opaque, triple lumen tube is introduced through the nose and allowed to migrate through the intestine by peristaltic motility until the leading tip reaches

the ileum (Figure 1). The location of the tube can be assessed through radiography. After test meal administration, ileal effluents are continuously collected for 8 h through one of the lumens. The infusion of a nonabsorbable marker (polyethylene glycol 4000, PEG-4000) into the intestine through another lumen of the tube allows determination of the total ileal effluent flow during this period. Combined with the administration of ^{15}N intrinsically labeled dietary protein, this method provides accurate estimates of nitrogen or AA digestibility at the ileal level and has been in use for over 25 y. A PubMed search for related studies was conducted with key terms of “ileal digestibility” OR “ileal protein digestibility” OR “ileal amino acid digestibility” AND “human” and the digestibility coefficients of animal and plant proteins processed by different methods are provided in **Supplementary Table 1**. The true ileal digestibility of AAs ranged from 91% (glycine) to 99% (tyrosine) in milk protein, and from 89% (threonine) to 97% (tyrosine) in soy protein (23). The true ileal AA digestibility of unlabeled whey protein and zein was recently reported with low values for zein, ranging from 24% for cysteine to 64% for glutamine (21).

Ileal digestibility of casein, whey protein, soy isolate, and concentrate, were also determined in ileostomates and were found to be high, varying from 97% to 100% (20).

As summarized in [Table 3](#), the procedure in ileostomates is relatively noninvasive, and is convenient for a crossover design; with the collection of total digesta, the use of nonabsorbable markers could be avoided. However, these patients might suffer from different gut disorders and their ileum will have morphological and microbiological modifications (47, 48); hence the measured digestibility values might differ from healthy volunteers. Additionally, the recruitment capacity of patients with a permanent ileostomy is limited. The naso-ileal intubation model is limited by the invasiveness of the procedure and a variable tolerance of the tube among subjects as well as interindividual variability for the tube migration. The placement of the tube could cause a concertina effect on the small intestine (49) which might confound the di- and tri-peptidase digestion and subsequent AA absorption, however, this needs to be confirmed. Use of nonabsorbable markers could introduce errors in measuring the ileal effluent flow rate (50). Lastly, the recycling of the isotope from test proteins in the endogenous protein leads to a minor underestimation (~1%) of ileal nitrogen digestibility (50). Moreover, neither model is suitable for routine digestibility measurements in humans, particularly in vulnerable age groups and with pathophysiological conditions. However, long-term experience in the implementation of this technique allows for the minimization of bias and the provision of accurate and repeatable values.

Indirect methods for measuring AA digestibility

Dual isotope tracer method.

A minimally invasive dual isotope tracer technique to measure true ileal AA digestibility has been recently developed to overcome the invasiveness and complexities around collecting ileal digesta in direct methods (37). In this method, 2 differently intrinsically labeled proteins, a test protein (^2H or ^{15}N) and a reference protein (^{13}C) of predetermined digestibility are simultaneously fed in a plateau feeding protocol. The ratio of plasma appearance of AAs from test to reference proteins at plateau with respect to the meal administered and corrected for AA digestibility of the reference protein allows determination of the true ileal AA digestibility of the test protein ([Figure 2](#)) (6, 37). Since the method measures plasma appearance of AAs to represent protein digestion and AA absorption throughout the small intestine, the term true AA digestibility (TAAD) will be used henceforth to differentiate from direct methods that measure digestibility at the terminal ileum. The method assumes that AAs from both test and reference proteins undergo similar first pass and splanchnic metabolism and enter a common pool after digestion and absorption; the ratio of test to standard AAs from this pool cancels out this uptake and metabolism. Another assumption of the method is the equivalent absorption of differently labeled AAs from the test and reference proteins, which is reasonable as stable isotopic

differentiations are not known (51, 52). An advantage of using the plateau feeding protocol is the attainment of a steady isotopic plasma enrichment which negates the effect of slow or fast response proteins in the test meal that could influence the rate of AA metabolism (53). This pattern of feeding is not habitual and AA digestibility could differ when administered as a bolus meal as is habitually consumed. However, the reserve capacity of pancreatic proteases is much higher (~10–15 times) than the volume required for protein digestion under normal physiological conditions, suggesting that the digestibility estimates might not drastically vary with different meal patterns (54, 55), although an effect on protein of low digestibility cannot be excluded and must be evaluated.

The viable options of reference proteins could be $\text{U-}^{13}\text{C}$ -labeled proteins that are commercially available and commonly consumed, such as spirulina protein ([Supplementary Table 1](#)), which is well digested and the constituent AAs have a stable isotopic carbon backbone (37). Although animal source proteins could also serve as good reference proteins, the production of intrinsically labeled casein or egg or meat protein in high quantities will be expensive. The digestibility of reference protein can be determined against a free AA mix (^2H or ^{15}N labeled) which is considered to be representative of a completely digested protein (37). The use of $\text{U-}^2\text{H}$ is preferred over an ^{15}N -labeled AA mixture, as the exchange of ^{15}N during transamination does not allow tracing back to the original carbon skeleton of the AA. Although a $\text{U-}^{13}\text{C}$ -free AA mixture can also be used as a reference, a protein versus protein comparison for the test and reference has been preferred since the rates of absorption, metabolism, and utilization of AAs derived from peptides are different from those of free AAs (56, 57). Here too, the label must be carefully chosen as the phenylalanine digestibility of ^{15}N -labeled spirulina was found to be 15% lower than $[\text{U-}^{13}\text{C}]$ -labeled spirulina (37, 58). Additionally, the TAAD of legume protein (mung bean) has been found to be similar with the use of either $\text{U-}^{13}\text{C}$ spirulina protein or $\text{U-}^{13}\text{C}$ free AAs as the reference, in healthy individuals (59) ([Table 4](#)) in a plateau feeding protocol, but this needs to be tested in a bolus feeding or a shorter repeated meal protocol. A mixture of free labeled AAs would also be a preferred choice for patients with impaired digestion and absorption functions such as those with cystic fibrosis, environmental enteropathy, and others (58, 60, 61). Another advantage is that the high digestibility of free AAs ensures a lower interindividual variability, as digestibility has an inverse association with variability, and thus provides more reliable reference values (62). The TAAD of $\text{U-}^{13}\text{C}$ spirulina had a mean IAA digestibility of ~85% (37) and this has been used as a correction factor while measuring the digestibility of different test proteins in the same population. This might introduce an error when measuring protein digestibility in a different group of individuals, as the mean variability of IAA digestibility in spirulina was found to be ~6% ranging from 3% for threonine and 12% for lysine (37). This could be addressed by measuring the reference protein digestibility in the same participant using a crossover design.

TABLE 3 Principle, strengths, and limitations of the AA digestibility and metabolic availability assays used in humans

Methods	Principle of measurement and equation	Strength	Limitation	Reference
Direct balance	<ul style="list-style-type: none"> Measures disappearance of ingested protein-derived AAs from intestinal lumen Calculated as the difference between the amount of ingested dietary AAs and that recovered in the terminal ileum <p>For intrinsically labeled test proteins: $TID = \frac{\text{Dietary AA intake} - \text{Ileal dietary AA flow}}{\text{Dietary AA intake}} \times 100$ </p> <p>For unlabeled test proteins: $TID = \frac{\text{Dietary AA intake} - (\text{Ileal AA flow} - \text{Endogenous AA flow})}{\text{Dietary AA intake}} \times 100$ </p> <ul style="list-style-type: none"> Two models are adopted in humans: naso-ileal intubation and ileostomy model 	<ul style="list-style-type: none"> Standard and direct method for measuring true ileal digestibility with repeatable results Provides digestibility estimates of all AAs in a single trial Ileostomy model is noninvasive and does not require a nonabsorbable marker to estimate flow rates of ileal effluents 	<ul style="list-style-type: none"> Naso-ileal intubation model is invasive, requires sampling of ileal digesta and not suitable for routine application in humans Uncertainties around the recovery of the nonabsorbable markers in intubation model Expensive assay Ileostomy model cannot be employed for healthy humans Errors associated with the measurement of endogenous AA losses if test proteins are not intrinsically labeled Overestimates AA availability of heat-treated and chemically processed foods Microbial colonization in terminal ileum of ileostomates could confound the digestibility estimates Discounts colonic absorption of AAs, if any Not applicable for routine measurements, vulnerable age groups, and pathological conditions with altered digestion and absorption 	(6, 19, 23, 48, 50, 106–108)
Dual isotope tracer	<ul style="list-style-type: none"> Measures appearance of ingested protein-derived AAs in systemic circulation Compares appearance of labeled AAs in plasma from intrinsically labeled food proteins to that of a simultaneously ingested but differently labeled reference protein of known digestibility in relation to the meal administered An example equation for ^2H-labeled test protein and ^{13}C-labeled reference protein: $TAAD = \frac{\text{plasma } ^2\text{H}_{AA\text{meal}} / ^2\text{H}_{AA}}{\text{plasma } ^{13}\text{C}_{AA\text{meal}} / ^{13}\text{C}_{AA}} \times 100 \times Dig_{SD} \times F_{ICF}$ 	<ul style="list-style-type: none"> Minimally invasive method requires blood collection Provides digestibility estimates of almost all AAs in a single trial Measures digestibility of proteins in habitually consumed meal preparations Suitable for application in humans and vulnerable groups 	<ul style="list-style-type: none"> Expensive, as requires test and reference proteins to be intrinsically labeled Indirect method. Digestibility of reference protein needs to be established in target population Uncertainty introduced by the transamination/deamination reactions Analytical complexity 	(25, 37, 59, 107)

(Continued)

TABLE 3 (Continued)

Methods	Principle of measurement and equation	Strength	Limitation	Reference
IAAO slope ratio	<ul style="list-style-type: none"> • Oxidation of a ^{13}C-labeled IAA (indicator) is used as a surrogate for protein synthesis • Compares oxidation response slope of an indicator IAA to graded intakes of test IAA from test protein to that of reference crystalline IAA • $MA = \frac{\text{slope of } ^{13}\text{C recovery for test protein}}{\text{slope of } ^{13}\text{C recovery for reference IAA}} \times 100$ • Bioavailability of crystalline IAA assumed to be 100% • Test IAA must be the first limiting IAA in the diet • Oxidation rates of indicator must be linear to the graded intakes of test IAA • IAAO slopes for test protein and reference IAA must have a common origin at a fixed base intake of test IAA in crystalline form 	<ul style="list-style-type: none"> • Noninvasive, requires breath collection and rate of CO_2 production • Analytically simple and relatively less expensive • Does not require test proteins to be intrinsically labeled, less expensive compared to other methods 	<ul style="list-style-type: none"> • Repeated measures design with higher subject burden, difficult for routine application in vulnerable groups • Provides MA estimate of 1 IAA at a time • The interindividual variability of the MA estimates is not reported 	(67, 70, 107)

AA, amino acid; Dig_{50d}, digestibility of the reference protein; F_{TGd}, transamination correction factor; IAA, indispensable amino acid; IAAO, indicator amino acid oxidation; MA, metabolic availability; TAAO, true amino acid digestibility; TID, true ileal digestibility.

A PubMed search was carried out to retrieve TAAO values of different proteins using keywords of “ileal AA digestibility” OR “ileal digestibility” AND “dual isotope tracer” AND “humans,” and is presented in Supplementary Table 1. High-quality proteins such as egg white, whole boiled egg, and chicken meat using the dual isotope method have reported digestibility estimates of 87%, 90%, and 92%, respectively (63), which were similar to digestibility estimates obtained by the direct ileal balance method (27, 28); however, plant protein digestibility values were lower than other studies (64, 65). This could be attributed to the differences in the test meal matrix between these experiments and type of processing of the test proteins (Supplementary Table 1). The interindividual variability of the digestibility estimates around the mean is reasonably small (overall CV <6%) when measured in apparently healthy humans. A major advantage of the dual isotope tracer method (Table 3) is enabling AA digestibility determination in habitually consumed diets in vulnerable populations of infants, children, pregnant women, and older adults, and in pathophysiological conditions.

Indicator AA oxidation slope ratio method.

This method was developed using the principles of the traditional slope ratio growth assay in which growth and feed efficiency (by carcass weight) in animals have been measured to estimate bioavailability of an IAA in a dietary protein (66). Although the slope ratio growth assay is an absolute standard of estimating IAA bioavailability, the requirement for several groups of animals for bioavailability assessment of a single IAA, prolonged adaptation to a test IAA intake, and invasiveness precludes its routine application in humans (66). The indicator amino acid oxidation (IAAO) slope ratio method is a noninvasive adaptation of the growth assay, in which oxidation of an orally administered ^{13}C -labeled “indicator” IAA is used as a proxy to assess the contribution of an unlabeled test dietary IAA to protein synthesis; the higher the oxidation of indicator IAA, the lower the protein synthesis (67). This method estimates the MA (includes digestibility, absorption, and utilization) of a single selected IAA in a dietary protein. It does so by comparing the oxidation response slopes of the labeled indicator IAA (usually 1- ^{13}C phenylalanine) to graded intakes of the selected test IAA from a test protein at subrequirement concentrations, to that of a reference crystalline IAA mixture (Figure 3). The IAAO slope ratio method has been validated against the growth assay with a mean difference of 4–7% for the MA estimates of lysine in differently processed pea proteins (68).

A key condition of the method is that the IAAO response must be linearly and inversely related to the test IAA intake concentrations, which requires the intakes to be at subrequirement concentrations, preferably <60% of the requirements (6). Due to the known high variability of IAA requirements, of ~25–40% (69), this critical condition of linearity could be challenging to satisfy, as has been observed previously, where the oxidation of indicator was reported

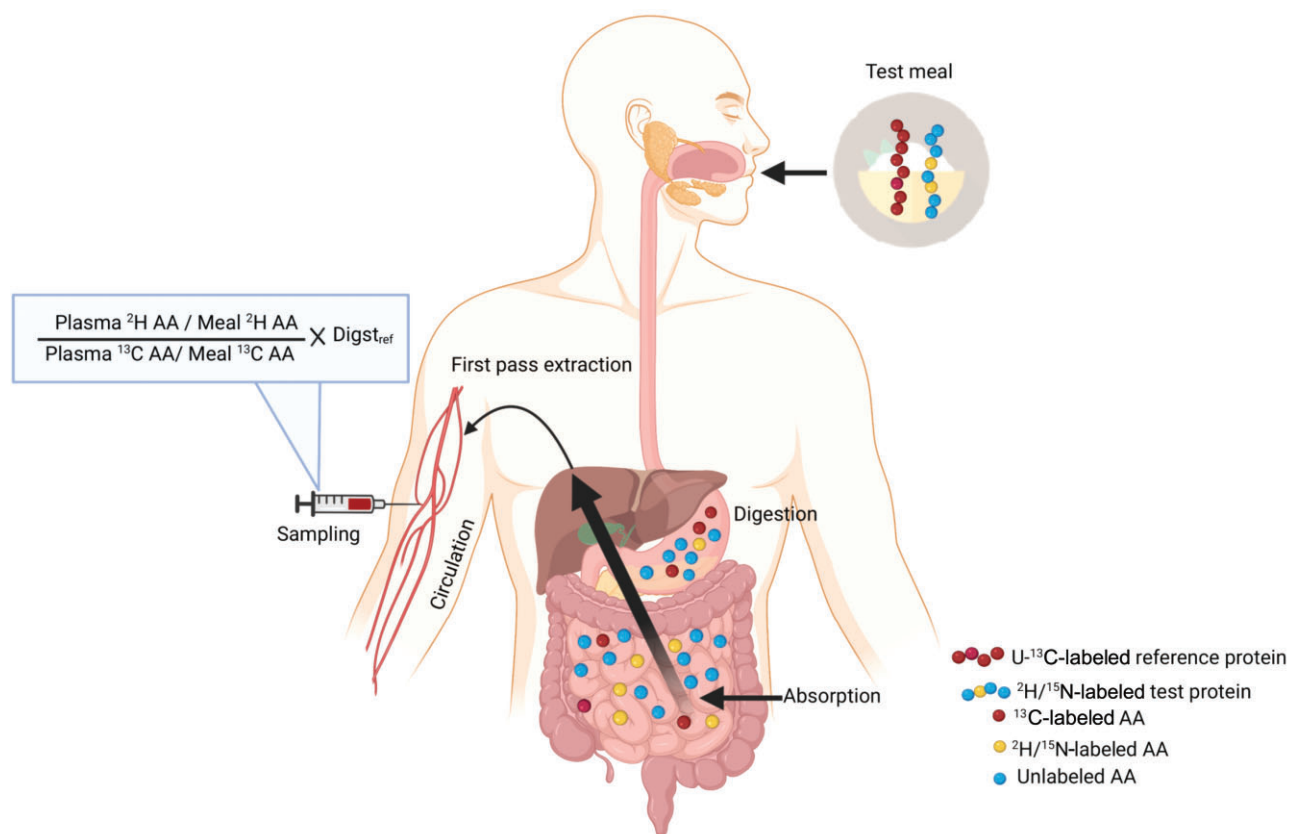


FIGURE 2 Schematic representation of the principle of the dual isotope tracer method to measure small intestinal AA digestibility. The test meal containing intrinsically $^2\text{H}/^{15}\text{N}$ -labeled test protein (yellow and blue circles) and uniformly ^{13}C -labeled reference protein (red circles) undergoes digestion. After absorption and first-pass splanchnic extraction, AAs from both the proteins enter the systemic circulation. The plasma appearance of individual AAs from $^2\text{H}/^{15}\text{N}$ -labeled test protein is compared with that of ^{13}C -labeled reference protein of known digestibility with respect to the test meal to determine the small intestinal AA digestibility in the test proteins. AA, amino acid; $\text{Digst}_{\text{ref}}$, digestibility of reference protein.

to plateau, when it should not have, at subrequirement concentrations of test IAA intakes (70). Further, it has been argued that a highly digestible protein (casein and hydrolyzed casein) could be a better control, as food-derived peptides have been shown to influence gut protein metabolism, stimulate gut endogenous AA secretions, and are better absorbed than free AAs (56, 57, 71).

An advantage of the IAAO slope ratio method is that it is noninvasive and analytically simple, involving only the measurement of $^{13}\text{CO}_2$ enrichment in noninvasive breath samples (Table 3). It does not require intrinsically labeled test proteins and is relatively cost-effective, particularly for mixed meals where all food protein sources in a mixed meal would need to be intrinsically labeled. The method provides reproducible results and is sensitive in the detection of postprocessing (heat treatment) reduction in MA as has been shown for lysine in peas, rice, and milk (68, 72–74). A PubMed search using key terms of “amino acid metabolic availability” OR “amino acid bioavailability” AND “indicator amino acid oxidation” AND “humans,” was performed to obtain MA of IAAs in different proteins, as measured in

humans, and the results are presented in Supplementary Table 1. The method does not report interindividual variability of the MA estimates, however, the CV appears to range from 15% to 52% across the studies when the SE of the slope was considered (70, 72, 73, 75). Nevertheless, the method provides reasonable MA estimates for foods and as expected, has been found to be lower than true ileal IAA digestibility or TAAD values (23, 43, 59, 72–74), except for rice, which could be due to the differences in milling processes (73, 75, 76).

The IAAO slope ratio method provides MA of a single IAA through multiple experiments with 3–4 graded intakes of test IAA in a repeated measures study design, which requires significant time, subject compliance, and meticulous control of their body composition, physiological status, and dietary intakes for the study duration. Moreover, the method provides a measure of metabolic utilization of a dietary protein. With rigorous application, the noninvasive IAAO slope ratio method has the potential to determine MA of limiting IAAs of dietary proteins in habitually consumed meal preparations in humans.

TABLE 4 True AA digestibility of mung bean with spirulina and crystalline AAs as reference proteins using dual isotope tracer method in healthy adults^{1,2}

True AA digestibility (%) IAA	MB ^{3,4}	MB- ¹³ C AA ^{3,4}
Methionine	52.2 ± 7.2	48.7 ± 6.3
Phenylalanine	73.4 ± 6.3	74.6 ± 1.4
Threonine	42.5 ± 1.2	42.7 ± 3.2
Lysine	63.0 ± 5.4	69.3 ± 3.4
Leucine	67.5 ± 3.2	69.3 ± 5.0
Iso-leucine	75.8 ± 2.6	76.6 ± 5.0
Valine	67.8 ± 6.0	66.7 ± 5.1
Mean IAA	63.2 ± 1.5	64.0 ± 2.4

¹Values are mean ± SD; AA, amino acid; IAA, indispensable amino acid; MB, mung bean true AA digestibility referenced to spirulina protein (*n* = 6); MB-¹³C AA, mung bean small intestinal IAA digestibility referenced to standard ¹³C IAA mixture (*n* = 5).

²Reproduced with permission from Kashyap et al. (59)

³Paired t-test between MB compared with MB-¹³C AA (*n* = 5), no significant differences were observed in IAA digestibility between the groups.

⁴No significant difference in true AA digestibility between MB compared with MB-¹³C AA.

Other methods of protein quality evaluation using stable isotopes.

In addition to the direct and indirect methods detailed above, isotopic methods allow the determination of postprandial utilization of dietary proteins and their contribution to protein synthesis. The net postprandial protein utilization (NPPU) can be determined using intrinsically ¹⁵N-labeled dietary proteins, based on the difference between total ingested dietary protein corrected by real/true ileal protein digestibility, and dietary nitrogen transfer to ammonia/urea by deamination (Supplementary Material 1). Using double labeled ¹⁵N-¹³C eggs, metabolic loss of AAs was ~18% as assessed by ¹³CO₂ and ¹⁵N recovery for an 8-h period (77). Nevertheless, there are almost no studies using ¹³C-labeled protein, due to the labeling cost as mentioned above. The NPPU estimates have been determined for different protein sources in adults with adequate protein adaptation and ranged from 72 to 78% for high-quality proteins, such as milk (78, 79) or soy protein (80), to 63–66% for lower quality proteins such as wheat (31).

The intravenous infusion of an isotopically labeled AA also allows the determination of dietary postprandial protein utilization (PPU) (81). In this method, leucine oxidation is measured in the postabsorptive and postprandial phases to determine leucine balance in relation to ingested leucine and a continuous infusion of 1-¹³C leucine (81, 82) (Supplementary Material 1). Dietary protein utilization was calculated by converting leucine balance to nitrogen balance in relation to nitrogen intake (82, 83). Depending on the feeding protocol, the PPU of wheat protein was 61–68%, which was consistent with NPPU estimates. However, the PPU of milk (93–100%) was found to be considerably higher than the NPPU estimates (79, 82, 84). The approach has not been used to determine digestibility of dietary proteins due to the theoretical assumptions and uncertainties associated with splanchnic sequestration of labeled AA, tracer recycling, issues with measurement of precursor enrichment, and the

assumed conversion factor of leucine to nitrogen content in tissue proteins (6).

Another method uses the combined ingestion of intrinsically labeled protein with a single AA (1-¹³C, ¹⁵N or ²H₅ phenylalanine, or 1-¹³C leucine) and an intravenous infusion of the same AA but differently labeled (²H₅ phenylalanine, or ²H₃ or 1-¹³C leucine) to measure endogenous and exogenous AA fluxes, and the contribution of exogenous AAs to whole-body and skeletal muscle protein synthesis (85). Postprandial protein handling and metabolism have been previously measured in high-quality animal and plant proteins (53, 86–91). The method can be used to assess the effect of protein source, protein load, exercise and meal timing on whole-body protein synthesis as well as muscle protein anabolism when associated with muscle biopsies (89, 92, 93). Although the technique could be used as a proxy for measuring protein quality, the use of a single intrinsically labeled test protein could underestimate the rate of plasma appearance of exogenous AAs due to the dilution and recycling of the tracer in the splanchnic bed and gastrointestinal tract (94). Additionally, in comparison to some of the other methods described in this review, it is invasive and not suitable for routine use.

Conclusion

The 2014 FAO expert consultation on protein quality evaluation recommended the use of DIAAS as the preferred metric for assessing the protein quality of individual foods and mixed diets (6). This requires the determination of individual AA digestibility at the ileal level which, until recently, was impossible to achieve without an invasive intubation or exteriorization of the intestine. However, together with the use of stable isotopes to intrinsically label dietary proteins, this is the only direct method that exists and has allowed accurate and precise digestibility estimates over the past 25 y and can be considered as a reference method for assessing protein and AA digestibility. Stable isotopes have been used as key tools in the field of protein and AA digestibility investigation and more largely protein quality, including in the development of minimal or noninvasive methods. The recently developed minimally invasive dual isotope tracer method can measure TAAD of almost all AAs in a single trial from dietary proteins in habitually consumed meal preparations in different age groups and shows promise to be applicable to vulnerable age groups and in those with pathophysiological conditions. However, the method is relatively new and needs to be validated against conventional assays which measure digestibility at the terminal ileum. The IAAO slope ratio method provides MA of one IAA at a time and demands significant time and compliance input from subjects in a crossover study design, which limits its use in vulnerable groups. Nevertheless, it is of interest due to its noninvasiveness and analytical simplicity.

Although efforts have been made to determine protein and AA digestibility of different foods, the agreement between the methods has not been rigorously evaluated by

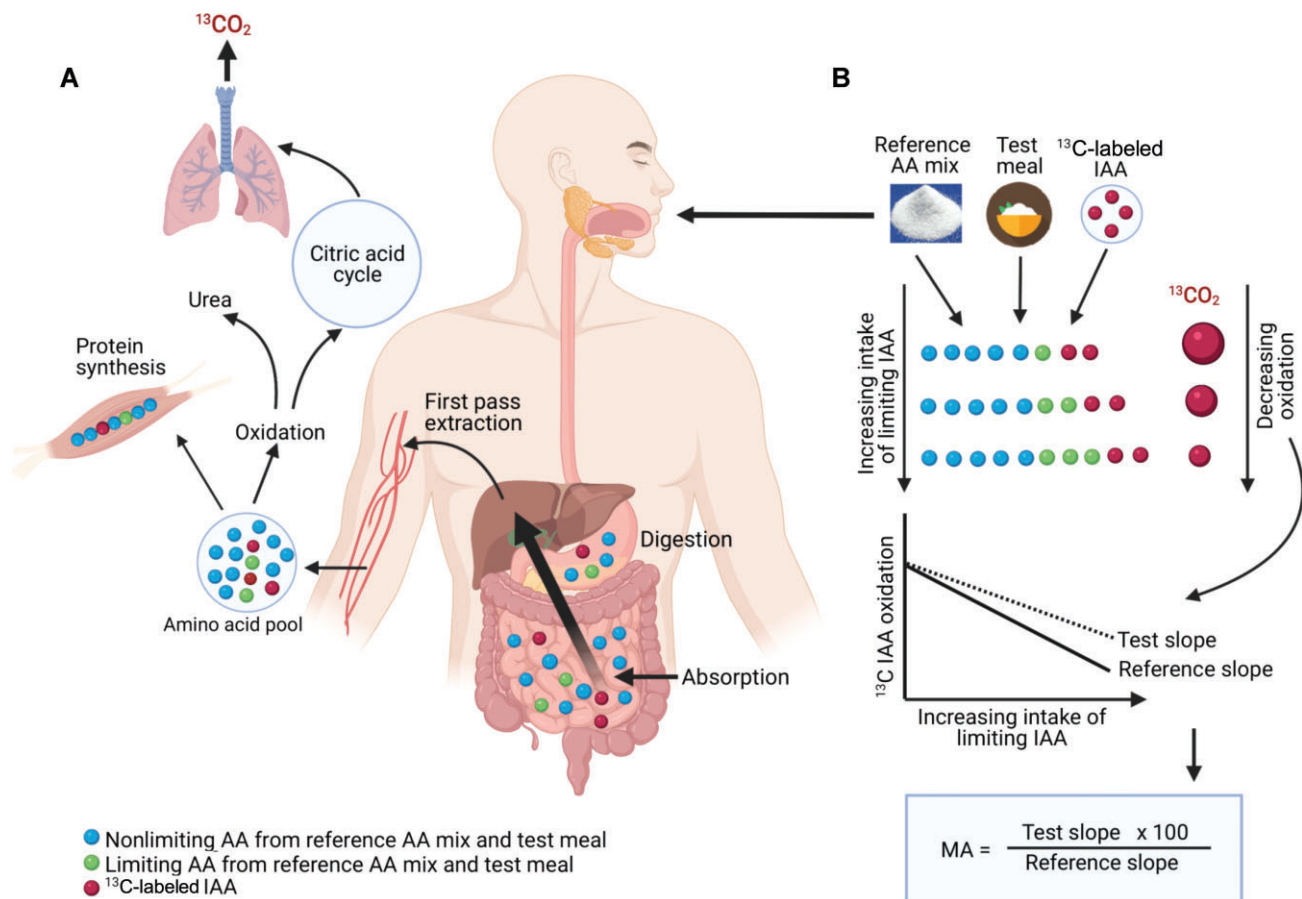


FIGURE 3 Schematic representation of the principle and application of the IAAO method to estimate MA of limiting AAs in test proteins. (A) Represents the digestion, absorption, and metabolic handling of free or protein-derived unlabeled nonlimiting (blue circles), limiting/test (green circles), and ^{13}C -labeled indicator IAA (red circles). (B) Subjects are provided with increasing intakes of limiting IAA (green circles) at the subrequirement concentration from a reference AA mixture or a combination of reference AA mixture and test protein with constant intake of ^{13}C -labeled indicator IAA (red circles) across the study days in a repeated measures design. With increasing intake of limiting/test IAA (green circles), the incorporation of ^{13}C -labeled indicator IAA (red circles) into tissue protein synthesis increases with the subsequent reduction in its oxidation, which is measured as $^{13}\text{CO}_2$ in breath. The IAAO response slopes are obtained by measuring both reference AA mixture and test protein at the same concentrations of limiting/test IAA intakes. The MA of limiting/test IAA in a test protein is computed by comparing the estimated IAAO response slope of test protein to the estimated slope of reference AA mixture, which is assumed to have 100% MA. AA, amino acid; IAA, indispensable amino acid; IAAO, indicator amino acid oxidation; MA, metabolic availability.

measuring the digestibility of the *same* protein source across methods. This is critical for harmonizing the digestibility values obtained from different methods considering the variable effect of the food matrix, of processing, and species-specific differences (particularly for plant proteins) on protein digestibility (95–97). As a start, the mean IAA and lysine digestibility of similarly processed whole-milk powders were found to be comparable across the methods (Supplementary Table 1), however, this needs to be expanded for other protein sources with low (<75%) and medium (<85%) digestibility. Until then, the values of true ileal protein and IAA digestibility/MA of different foods that are presented in Supplementary Table 1, could serve as a

guide for field-level use. Further studies are also required to establish the acceptable limits of interlaboratory variability in true ileal protein and IAA digestibility/MA measurements of foods.

With the development of DIAAS, the re-evaluation and harmonization of these methods is important for supporting protein content claims of food labeling systems across countries. Moreover, efforts to determine the protein and AA digestibility in relevant food groups and combinations using minimally invasive methods need to be continued to build into the expanding database of DIAAS and to inform the agricultural supplementary nutrition programs and industrial regulatory frameworks and policies.

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