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# Determinants of amino acid bioavailability from ingested protein in relation to gut health

Claire Gaudichon and Juliane Calvez

## Purpose of review

The current review provides an update on the recent research developments regarding amino acid bioavailability in conditions of both good health and gut disorders.

## Recent findings

Determination of amino acid bioavailability is complex and invasive. Minimally invasive methods using stable isotopes have been developed for humans. Data were collected in different models – humans, pigs and rats with various procedures – leading to interstudy variability. They mainly focused on either plant protein or the effect of food processing on animal protein. Plant protein in their original food matrix (legumes, grains, nuts) are generally less digestible (about 80%) than animal protein (meat, egg, milk; about 93%). Food processing has a limited impact on animal protein but its effect might be higher on plant protein. Few studies have documented the effect of gut disorders on protein digestibility, except in gastric bypass where paradoxical effects were reported. Data are needed to identify the amplitude of protein malabsorption in diseases such as inflammatory bowel disease or environmental enteric dysfunction.

## Summary

The past 5 years have seen a renewed interest in amino acid bioavailability in view of assessing protein quality to support current shifts in protein sourcing. Methodological developments have been performed and several studies have reported values in various models. The question of protein digestibility in gut disorders remains poorly addressed.

## Keywords

animal models, digestibility, humans, plant and animal proteins, stable isotopes

## INTRODUCTION

Amino acid bioavailability can be defined as the proportion of amino acids reaching systemic circulation and that can be incorporated into body protein synthesis. It mainly depends on the protein digestibility and amino acid absorption as well as the latter's metabolic fate in deamination pathways. Although metabolic losses are a component of amino acid bioavailability, most studies refer to the efficiency of protein digestion. The term 'protein and amino acid digestibility' generally refers to the amount of these undigested components recovered in the digesta. The term 'amino acid bioavailability' is related to absorbed amino acids. However, these terms are not fully consistent in the literature. In the context of designing strategies to ensure protein security in the future while responding to environmental issues [1], there has been a renewed interest in protein digestibility as a key factor of protein quality in the 5 past years. Many studies were conducted in healthy volunteers or their related animal models. Whereas digestive disorders can compromise protein security in some populations due to protein

malabsorption, little objective data exist. This review aims to provide a state of the art of recent developments related to protein and amino acid digestibility in normal and abnormal gut function.

## METHODOLOGICAL CONSIDERATIONS FOR MEASURING AMINO ACID BIOAVAILABILITY

Methods for determining digestive efficiency, namely digestibility, have evolved in the context

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## KEY POINTS

- Amino acid digestibility must be measured at the ileal level or using alternative procedures developed recently, in human and animal models.
- The amino acid bioavailability of legumes, refined and brown cereals, nuts, eggs, milk and meat with various treatments has recently been determined under various methodological conditions.
- The interindividual variability is inversely associated with digestibility and can be assumed as a marker of low amino acid bioavailability.
- In the case of gastric bypass, protein digestive efficiency is improved due to intestine hypertrophy, but metabolic bioavailability is paradoxically decreased.
- Limited data exist for other unhealthy gut conditions, although protein malabsorption has been quantified in pancreatic insufficiency and short bowel syndrome.

of the Food and Agriculture Organization of the United Nations-recommended Digestible Indispensable Amino Acid Score (DIAAS) values as the preferred criteria of protein quality [2]. In this context, digestibility of each indispensable amino acid (IAA) must be measured at the ileal level because absorption occurs in the small intestine. Amino acids reaching the large intestine are fermented, and molecules are partly transformed into either other amino acids or metabolites that can be absorbed at the colon level, such as ammonia or indoles for nitrogen molecules, or H<sub>2</sub>S, phenol and branched-chain fatty acids for nonnitrogenous compounds [3]. If the measurement of protein digestibility at the faecal level has been considered an overestimated but acceptable proxy of ileal digestibility, the measurement of faecal amino acid digestibility is generally not accepted, though already reported for pulse proteins, for instance [4].

It must be mentioned that the term 'amino acid availability' is sometimes used on the basis of the plasma amino acid area under the curve (AUC) after a test meal, but it is confusing because it does not reflect amino acid absorption alone [5,6]. Indeed, other phenomena such as the speed of digestion and the efficiency of amino acid uptake in tissues considerably influences the incremental plasma amino acid postprandially. As an illustration of the confusion that can occur, casein generally displays a moderate AUC compared with other protein sources, such as whey; a result that is not ascribed to a low amino acid bioavailability from casein since it is one of the most digestible proteins, but rather to a slow digestion rate. As the entry of amino acids into

protein synthesis rapidly saturates the process, there are more circulating, remnant amino acids when proteins are digested rapidly than when amino acids arrive progressively in the blood. When dietary protein is intrinsically labelled with a stable isotope and a labelled amino acid is intravenously infused, it is possible to determine the amount of dietary amino acids that appear in circulation [7]. However, owing to splanchnic extraction, this value reflects the availability of dietary amino acids for peripheral organs but not digestibility.

Measuring ileal digestibility is complex and requires invasive procedures, depending on the model. In humans, intestinal tubing allows for the continuous collection of effluents and adequate measurements of ileal amino acid losses [8<sup>¶</sup>]. However, this method is very invasive and ileostomized patients have therefore been used as convenient alternative due to the direct access to digesta. Nevertheless, there are currently few eligible patients, making recruitment difficult. Recently, a minimally invasive method has been developed in healthy volunteers. The dual isotope method is indirect as it is based on the use of a reference protein in the meal together with the test protein. The latter is labelled with one isotope, preferably <sup>2</sup>H [9<sup>¶</sup>], or alternately <sup>15</sup>N [10] and the reference protein (spirulina or free amino acids) with <sup>13</sup>C. The amino acid absorption from the test protein is determined by analyzing the relative ratio of these two tracers in the meal and the plasma amino acids. The low invasiveness is a major strength of this method, but it is analytically complicated. Moreover, it involves indirect calculations and to date, it has not yet been validated against the direct method of ileal sampling.

An entirely noninvasive method has been developed from the indicator amino acid oxidation (IAAO) method that was primarily used to determine amino acid requirements. It was adapted to identify the metabolic availability (i.e. including digestive and metabolic losses) of the limiting amino acid [11<sup>¶</sup>]. The principle is that if one amino acid is limiting in the diet, the other amino acids are subjected to increased oxidation because their entry into protein synthesis is impaired. This is thus reflected by the increased <sup>13</sup>CO<sub>2</sub> excretion in breath air from the indicator amino acid, generally <sup>13</sup>C phenylalanine. The method consists in giving either synthetic meals with increasing levels of the limiting amino acid in a free form (thus theoretically 100% digestible) or meals containing the test protein with the same levels of the limiting amino acid. The amino acids are given under the requirement level to obtain a linear regression between the amino acid level and the <sup>13</sup>CO<sub>2</sub> excretion. The slopes obtained with free amino acids and the protein

meals are then compared, reflecting a lower bioavailability of the limiting amino acid in the test protein if the slope is weaker than with the free amino acid meal. As with the dual isotope method, it is an indirect method. It has the advantage of being analytically simple and entirely noninvasive, but it is cumbersome to formulate the meals and then adapt volunteers to different ones. Moreover, it ultimately only measures the bioavailability of the limiting amino acid.

Animal models are also used to provide amino acid digestibility data, the pig model equipped with an ileal or ileocaecal cannula that allows the continuous recovery of digesta, in particular [12]. However, ethical concerns can limit the use of this model. Rats offer an alternative model for digestion studies. They are much less expensive and easy to use, but a major drawback is the impossibility of continuously collecting the ileal digesta. To circumvent the problem, two strategies exist. One consists of collecting the ileal digesta at a single time postmeal, with a repeated meal protocol [13<sup>■</sup>]. The second strategy is to collect all the digesta in the caecum 5–6 h after meal ingestion, a period of time that allows complete meal digestion but limits the fermentation processes [14<sup>■</sup>]. Both methods present some limitations but allow discrimination of protein digestibility in different conditions. Lastly, another methodological factor of variation is the method of differentiation between dietary and endogenous protein losses by either using a protein-free diet to assess the endogenous losses or stable isotopes to trace dietary or endogenous proteins [3].

### DETERMINANTS OF AMINO ACID BIOAVAILABILITY FROM INGESTED PROTEIN IN HEALTHY GUT CONDITIONS

In normal physiological conditions, when digestive capacities are not impaired, the main determinant of digestive bioavailability is the nature of the protein as well as the food matrix in which it is incorporated – protein isolates being generally more digestible than proteins in their crude matrices. Table 1 presents a representative set of the recent values of protein/amino acid digestibility, obtained in different models and with different methodologies described above. Both are major factors of variation that must be considered for any data comparison from the literature.

Notably, the bioavailability of amino acids from plant sources has been substantially reported owing to the increasing interest in evaluating the quality of nonconventional protein sources. Data obtained in humans with the dual isotope method indicate a mean IAA bioavailability in three different legumes

of 63–74% [16], whereas the value is around 90% for meat and eggs [17]. However, the amino acid digestibility of whole eggs can be severely decreased, down to 75%, when ingested with black tea [18], a result that can be ascribed to the polyphenols. Using ileal tubes to measure the true ileal digestibility (endogenous protein being evaluated in a protein-free group), mean amino acid bioavailability from whey was close to that of egg and meat, whereas that of zein was very low [8<sup>■</sup>], a phenomenon that can be partly ascribed to the isolate's low solubility. Using the IAAO method, Rafii *et al.* [19] determined the metabolic availability of the limiting amino acids in corn meal, lysine and tryptophan, and found higher values for these individual amino acids than those found for zein by Calvez *et al.* [8<sup>■</sup>]. It must be pointed out that in the latter study, the digestibility of lysine and tryptophan could not be determined because zein was almost completely deficient in both amino acids. Rafii *et al.* [11<sup>■</sup>] also reported that the most limiting amino acid in chickpea, methionine, was only bioavailable at 63% against 100% in steamed rice. In rats, our team found a high digestibility of a sunflower isolate, almost as high as that observed for whey in the same model, whereas the digestibility of spirulina cell amino acids was moderate [20]. In rats, the amino acid digestibility from different cereals was shown to be higher in wheat than in rice, and especially in polished rice [13<sup>■</sup>].

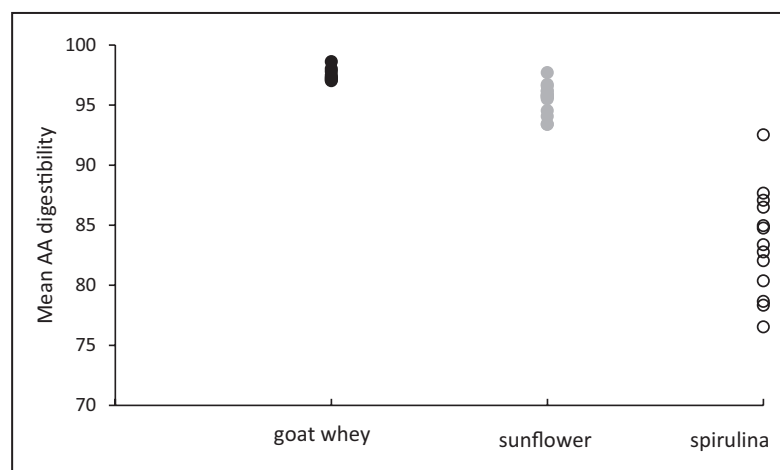
The treatments applied to food are also a determinant of bioavailability. The effect of bovine meat cooking processes such as boiling and grilling were reported to be modest in pigs, with differences of 2.5% between grilled and boiled beef [21]. This is in accordance with a previous result in rats, reporting the same amplitude of variation between roasting, grilling, barbecuing and boiling, except that the lowest digestibility was observed for boiled meat [22], in contrast to the study cited above. In a pig model, it was also observed that the cooking temperature of pork products, varying from 63 to 72 °C, did not modify amino acid digestibility; neither did the smoking process applied to bacon, however, curing improved the amino acid digestibility of ham from 95 to 99% [23<sup>■</sup>]. In contrast, heat treatment applied to nuts had an inverse effect with a decrease of pistachio protein digestibility by 10% after roasting, as measured in pigs [24].

Significantly, an inverse relationship between digestibility and interindividual variability has consistently been observed. In other words, when a protein is highly digestible, all individuals display a great capacity to digest the protein, but when the protein is less digestible some of them have a great capacity to digest it while others have a low capacity. This is illustrated in Fig. 1, which depicts the

**Table 1.** Amino acid digestibility of various protein sources in recent studies using different models and methodologies

Protein source	Parameter	Bioavailability (%)	Model	Method <sup>a</sup>	Reference
Whey	Mean AA	92 ± 6	Humans	Nasoileal tube/protein-free group	Calvez <i>et al.</i> [8 <sup>■</sup> ]
Zein	Protein	91 ± 6			
	Mean AA	63 ± 13			
	Protein	60 ± 13			
Chickpea	Mean IAA	74.5 ± 0.8	Humans	Dual isotope method/repeated meals	Kashyap <i>et al.</i> [16]
Mungo beans	Mean IAA	63 ± 1.5			
Yellow pea	Mean IAA	71.5 ± 1.5			
Egg	Mean IAA	89.5 ± 4.5	Humans	Dual isotope method/repeated meals	Kashyap <i>et al.</i> [17]
Meat	Mean IAA	92 ± 3			
Chickpea	Methionine	63	Humans	IAAO	Rafii <i>et al.</i> [11 <sup>■</sup> ]
Rice	Methionine	100			
Corn meal	Tryptophan	80	Humans	IAAO	Rafii <i>et al.</i> [19]
	Lys	71			
Meat	Mean AA	96.5 (grilled) to 98.5 (boiled)	Pigs	Single ileal sample/repeated meals/endogenous losses from the literature	Hodgkinson <i>et al.</i> [21]
Pork products	Mean AA	95 (Noncured ham) to 100 (raw belly) ± 2.5	Pigs	Ileal cannula/protein-free diet	Bailey <i>et al.</i> [23 <sup>■</sup> ]
Pistachio	Mean AA	85 (Roasted) to 95 (raw) ± 7	Pigs	Ileal cannula/protein-free diet	Bailey and Stein [24]
Sunflower isolate	Mean AA	96 ± 0.5	Rats	Postprandial caecal-faecal losses/intrinsic <sup>15</sup> N labelling	Tessier <i>et al.</i> [14 <sup>■</sup> ]
Goat whey	Mean AA	97.5 ± 0.5			
Pea isolate	Mean AA	94.5 ± 4	Rats	Postprandial caecal-faecal losses/protein-free group	Guillin <i>et al.</i> [15]
Gluten	Mean AA	94.5 ± 3.5			
Spirulina whole cells	Mean AA	83.5 ± 4.5	Rats	Postprandial caecal-faecal losses/intrinsic <sup>15</sup> N labelling	Tessier <i>et al.</i> [20]
Brown rice	Mean AA	85 ± 2.3	Rats	Single ileal sample/repeated meals/endogenous losses from the literature	Han <i>et al.</i> [13 <sup>■</sup> ]
Polished rice		77 ± 8.5			
Buckwheat		88 ± 6			
Whole wheat		93 ± 2			

Bioavailability values are presented as mean ± SD. AA, amino acid; IAA, indispensable amino acid; IAAO, indicator amino acid oxidation.  
<sup>a</sup>Type of method, sample collection, specific feeding procedure, procedure to determine true digestibility (assessment of endogenous intestinal losses) or real digestibility (stable isotopes to trace dietary protein).



**FIGURE 1.** Interindividual dispersion of amino acid digestibility in rats depending on the protein source. Rats were given a <sup>15</sup>N test meal and <sup>15</sup>N amino acid losses were determined in the caecum 6 h after the ingestion to determine the oro-caecal amino acid digestibility. <sup>15</sup>N goat whey: n = 8; <sup>15</sup>N sunflower isolate: n = 15; <sup>15</sup>N spirulina (whole cell): n = 14. Adapted from [14<sup>■</sup>,20].



individual data of amino acid bioavailability in rats for  $^{15}\text{N}$  goat whey, sunflower isolate and spirulina. These three data sets are directly comparable because they are methodologically uniform. It appears that for spirulina, which has a digestibility 12–15% lower than that of the other two proteins, the data dispersion is dramatically expanded. We observed the same in humans, for instance, concerning zein digestibility, which was the lowest we observed among our intestinal tube studies [8<sup>■</sup>]. The SD was 13% (for a mean of 60%), with the lowest individual value being 39% and the highest 82%. This is consistent with a previous observation of rapeseed isolate that contains resistant albumin (napin), with a mean value of  $84 \pm 8.8\%$  and extreme values of 65 and 90%, as reviewed recently by Walther *et al.* [25]. In this review, the authors proposed variations in protease activity and polymorphism of amino acid transporters as putative causes of interindividual variability, but data are lacking to support these hypotheses. The microbiome may also play a role as indirectly suggested by an intervention study in humans where the administration of vancomycin increased faecal energy losses [26]. However, the rationale for a putative role of the large intestine microbiota in amino acid bioavailability is unclear since there is no significant absorption of amino acids in the colon.

## PROTEIN DIGESTIBILITY IN DIGESTIVE DISORDERS

In digestive disorders there are little data on protein and amino acid bioavailability, especially in recent years. Protein malabsorption has been observed or supposed in different situations such as bariatric surgery, short bowel syndrome (SBS), pancreatic insufficiency, inflammatory bowel disease (IBD) and environmental enteric dysfunctions (EED).

In the 3 last years, the most documented unhealthy gut condition is related to bariatric surgery. Indeed, after a gastric bypass, which drastically modifies the gut anatomy, amino acid bioavailability may be altered in a paradoxical manner. Significantly, the blood appearance of amino acids from  $^{13}\text{C}$  labelled casein was shown to occur faster after gastric bypass than in normal volunteers [27<sup>■</sup>]. This result was also associated with a faster glucose handling by the intestine. It is thus likely that bioavailability of amino acids from ingested protein is not impaired by a gastric bypass. Our team confirmed this hypothesis in a gastric bypass rat model in which we observed that  $^{15}\text{N}$  labelled milk protein was paradoxically more digestible in operated rats than in sham rats [28]. This better digestive

efficiency was slight (+2%) but significant. In this study, we also evaluated the metabolic bioavailability of dietary nitrogen (N) by measuring the  $^{15}\text{N}$  retention in tissues. We found a lower dietary N retention in organs, especially liver and muscle, a paradox that can be explained by a higher retention of dietary N in intestinal mucosa. Indeed, as previously observed, the mucosa was hypertrophied and the  $^{15}\text{N}$  retention/g of tissue was as high in gastric bypass as in sham rats. Gastric bypass but not sleeve gastrectomy slightly improves digestive amino acid bioavailability but lowers metabolic bioavailability which could thus contribute to protein deficiency [29]. Energy absorption was evaluated in children with three types of SBS, differing by the length of the bowel and the presence or not of the colon and ileocaecal valve [30<sup>■</sup>]. The energy absorption was only 60–68% and was linearly linked to the length of the small bowel, with a better efficiency in the presence of a remnant colon. The patient characteristic data indicated that apparent protein digestibility was 57–65%, without any effect of the type of SBS, suggesting a protein malabsorption of about 20–30% in this syndrome.

In gut disorders involving pancreatitis insufficiency, protein malabsorption has been little documented in contrast to lipids, of which absorption is majorly impaired. After ligation of the pancreatic duct in minipigs, we observed a drastic alteration of real ileal protein digestibility that was  $29 \pm 11$  vs.  $89 \pm 6\%$  in control pigs [31]. The administration of pancreatic enzymes partially restored protein absorption in a dose-dependent manner. Significantly, the ligation of the canal duct did not completely suppress protein digestion as about 30% of dietary protein was digested. This indicates that nonpancreatic enzymes, such as pepsin and/or brush border enzymes, can partially compensate for the experiment-induced pancreatitis insufficiency. Cystic fibrosis (CF) is also associated with pancreatic insufficiency. Using the principle of the dual isotope method described above, Engelen *et al.* [32] reported that the digestibility of phenylalanine from spirulina was 80% in healthy volunteers but less than half in CF patients. This is a dysfunction that could be counteracted by enzyme substitution, as reported for chronic pancreatitis.

In inflammatory gut disorders, such as EED and IBD, there are almost no data on protein digestibility. Protein malabsorption can be suspected due to diarrhoea, intestinal mucosal abrasion and alteration of permeability, leading to undernutrition, especially in children [33]. Devi *et al.* [34<sup>■</sup>] indirectly addressed this question by assessing amino acid bioavailability in moderately stunted Indian children. They used the ratio kynurenine/tryptophan as a

probable biomarker for intestinal inflammation. This marker was negatively associated with height-for-age Z score. The authors observed a positive association between the kynurenine/tryptophan ratio and proline digestibility, which they interpreted as a reflection of the specific role of metalloproteinases from the brush border membrane that both respond to inflammatory stress and activate proline-rich peptide cleavage. However, they did not find any correlation with the digestibility of other amino acids. In addition, the mean amino acid bioavailability of chickpea ( $87 \pm 10\%$ ) and yellow pea ( $86 \pm 10\%$ ) were higher than what was found in adults, as reported in the former paragraph [16]. It is noteworthy, though, that the interindividual variability is greater and thus suggests an impairment of amino acid bioavailability. However, the experimental conditions, without any direct evaluation of gut dysfunction or inflammation, do not permit drawing conclusions regarding the absence of any protein absorption alteration in EED.

## CONCLUSION

The past few years have seen a renewed interest in protein and amino acid bioavailability in view of documenting the quality of plant and alternative proteins to animal proteins and, furthermore, to enrich the DIAAS database. As protein and amino acid digestibility are methodologically complex and invasive, many developments have been conducted to reduce the invasiveness or to improve the accuracy of data. In conditions of healthy gut functioning, the true digestibility depends on the protein source, animal more than plant protein, especially in the context of the food matrix where several values are below 75% for plants. Important interindividual variability is observed for low digestible protein but the reasons remain unknown. In abnormal gut functions, protein digestibility has been little-addressed and dedicated studies to determine protein malabsorption and the subsequent risk of protein malnutrition are required.

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## Conflicts of interest

There are no conflicts of interest.

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- of outstanding interest

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