

Comparative Nutritional Value for Amino Acids, Oligopeptides and Soybean Protein

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In recent years our view on the absorption of protein digestion products has undergone a radical change. We now believe that intralumen hydrolysis of intake proteins to free amino acids is only partial. It has thus been hypothesized that partial protein hydrolysates or oligopeptide mixtures are nutritionally superior to corresponding amino acid mixtures. With this consideration for a background, we developed a sophisticated technique of protease-catalyzed modification of soy protein to produce enzymatically modified proteins (EMP) having different levels of covalently attached methionine. We also carried out feeding tests with protein-malnourished rats, with the result that for their recovery from a state of malnutrition, an EMP containing peptide-bound methionine at 3% was significantly more effective than the corresponding amino acid mixture and even the soy protein itself. Discussions stress the importance of applying such a technique to production of an oligopeptide rather than a free amino acid nitrogen source as a foodstuff for therapeutic use.

Recent research on protein nutrition has elucidated the digestion and absorption mechanism involving peptides. It is now believed that dipeptides and tripeptides formed by intralumen hydrolysis of intake proteins are mostly absorbed prior to further hydrolysis to free amino acids (1-5). Presumably, therefore, supplementation of low-quality proteins is more effective when essential amino acids are used as peptides rather than as free amino acids (6,7). To test this hypothesis, we must examine differences in digestion and absorption features, protein efficiency ratio (PER), and many other nutritional indices between an oligopeptide mixture (OPM) and the corresponding free amino acid mixture (AAM).

In this study we undertook experiments to examine nutritional effects of supplemental peptide-bound methionine with a soy protein isolate (SPI) and those of OPM products that may be usable as alimentary diets for recovery from a state of protein malnutrition.

EXPERIMENTAL

Preparation and characterization of enzymatically modified proteins. To prepare the peptide-bound methionine we utilized a papain-catalyzed reverse reaction which could proceed under an unusual condition (8-16), and then we compared the resulting enzymatically modified protein (EMP) with the AAM of the same amino acid

composition. A recrystallized preparation of papain (EC 3.4.22.2) was used to modify SPI with covalent incorporation of L-methionine. In this case, methionine was used in ethyl ester form (8). The conditions for the modification were set according to the previous paper (10). These were as follows: medium, 1 M carbo-

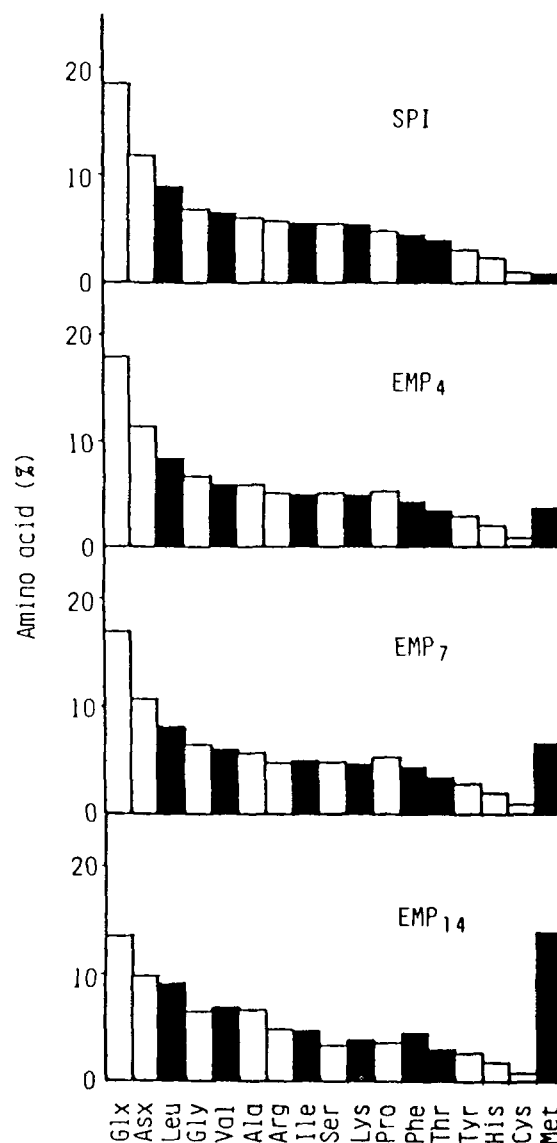


FIG. 1. Amino acid patterns of SPI and a series of EMP products. Suffixed figures stand for approximate percentage of methionine contents of the EMP products. Filled columns indicate essential amino acids.

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nate (pH 9) containing 10 mm L-cysteine; concentration of SPI in the medium, 25%; concentration of papain in the medium, 0.25%; incubation temperature, 37 C, and incubation time, 2 hr. After the incubation the mixture was dialyzed against water and an EMP product was obtained by lyophilization of the non-diffusible fraction. The methionine content of the EMP was adjusted by controlling the amount of methionine ethyl ester added to the reaction medium. Thus, we obtained a series of EMP products whose methionine levels were altered progressively (Fig. 1). Chemical analyses showed that every EMP product was a peptide mixture having a molecular weight distribution in the range of 2,000–12,000 daltons with an average at approximately 4,000 daltons. The positions of the methionine molecules incorporated were estimated by treatment of EMP₇ (Fig. 1) with exopeptidases. The treatment with leucine aminopeptidase indicated that no significant amount of methionine residues existed at or near the N-terminal position. On the other hand, by treating this protein with carboxypeptidase A, we found that the methionine residues were mostly located at or near its C-terminal position (17). From these results and other chemical analyses it was estimated that these methionine molecules existed in the form of an oligomer at the C-terminal position.

Digestion and absorption studies with EMP. Feeding tests were carried out with seven-week-old male rats (Wistar strain) weighing approximately 200 g each which had been imprinted for meal feeding. Two kinds of diets were prepared in accordance with Harper's

TABLE 1

Diet Composition

Ingredient	Composition (weight percent)	
	EMP ₃ diet	AAM diet
EMP ₃	10.5 ^a	—
AAM	—	9.7 ^a
Corn Starch	69.15	69.95
Sucrose	10.0	10.0
Soybean oil	5.0	5.0
Mineral mixture ^b	4.0	4.0
Cellulose powder	1.0	1.0
Choline chloride	0.15	0.15
Vitamin mixture ^c	0.20	0.20

^aEquivalent to 1.5% nitrogen.

^bAmounts of salts in 100 g of the mineral mixture: CaCO₃, 29.29 g; CaHPO₄•2H₂O, 430 mg; KH₂PO₄, 34.31 g; NaCl, 25.06 g; MgSO₄•7H₂O, 9.98 g; Fe(C₆H₅O₇)•6H₂O, 623 mg; CuSO₄, 156 mg; MnSO₄•H₂O, 121 mg; ZnCl₂, 20 mg; KI, 0.5 mg; (NH₄)₆Mo₇O₂₄•4H₂O, 2.5 mg; and Na₂SeO₃•5H₂O, 1.5 mg.

^cAmounts or units of vitamins in 100 g of the vitamin mixture: thiamine hydrochloride, 100 mg; riboflavin, 100 mg; niacinamide, 500 mg; calcium pantothenate, 400 mg; pyridoxine hydrochloride, 100 mg; folic acid, 10 mg; menadione, 10 mg; d-biotin, 4 mg; vitamin B₁₂, 0.6 mg; ascorbic acid, 1 g; vitamin E acetate, 8 g; vitamin A acetate, 80,000 units; and vitamin D₂, 8,000 units.

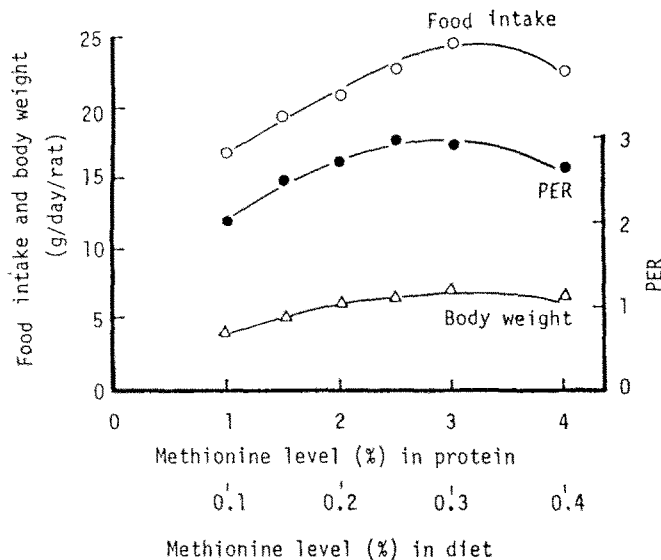


FIG. 2. Food intakes, weight gains and PER values observed for rats fed the SPI-EMP₁₀ formulas having different methionine levels.

composition (18); one was based on an EMP containing methionine at a 3% level (EMP₃) and the other on the simulating AAM formulated similarly. Each of the diets was also set at a nitrogen level of 1.5% (Table 1). The diets were administered to rats by meal feeding per os, and portal blood samples were taken every hour thereafter. At the same time, the contents of their small intestinal tracts were washed out with water and, immediately, treated with 10% trichloroacetic acid. Subsequently, the resulting supernatant was taken and fractionated on the basis of molecular weight by membrane filtration to obtain fractions of lower than 500 daltons, 500–1,000 daltons and 1,000–2,000 daltons. We also carried out another experiment to enhance the nutritional value of SPI by adding EMP₁₀, which stands for an EMP containing methionine at a 10% level. Diets were prepared according to Harper's composition (18) with seven different methionine levels varying in the range of 0.1–0.4% (Fig. 2), where nitrogen levels were all adjusted at 1.5%. Each of the diets was fed ad libitum to four-week-old male rats (Wistar strain) weighing approximately 80 g each. After the feeding of the rats was conducted for four weeks, their food intakes and body weight gains were measured to obtain PER values.

Preparation and characterization of oligopeptide mixture (OPM). We prepared an OPM product by hydrolyzing EMP₃ with trypsin. EMP₃ (100 g) was dissolved in water (5 l) and adjusted to pH 7.8 with NaOH. The solution was incubated with a recrystallized preparation of trypsin (100 mg) at 37 C for 15 min. Absolute ethanol (10 l) was added to stop the enzymatic reaction and the resulting 66.6% ethanol solution allowed to stand for 6 hr at room temperature. A precipitate formed and was removed by centrifugation at 4,000 rpm for 10 min; the supernatant was saved. The precipitate (42 g on a dry-matter basis) was submitted to a repetition of

TABLE 2

Amino Acid Patterns of SPI and OPM₃

Amino acid	SPI (g amino acid/16 g nitrogen)	OPM ₃
Asx	11.13	12.12
Thr	3.65	3.69
Ser	4.81	4.91
Glx	20.42	21.44
Pro	5.66	2.43
Gly	3.97	4.45
Ala	4.33	4.62
Cys	1.04	1.10
Val	5.04	5.20
Met	1.07	3.11
Ile	4.78	5.52
Leu	7.05	7.59
Tyr	3.79	3.57
Phe	5.25	5.92
Trp	0.98	0.99
Lys	5.72	5.42
His	2.65	2.02
Arg	7.22	5.52

the tryptic hydrolysis followed by the ethanol treatment to obtain a second supernatant. The first and the second supernatants were pooled prior to evaporation of the ethanol in vacuo. Lyophilization of the residue gave an OPM product in a yield of 87 g. Amino acid analysis and gel chromatography demonstrated that the OPM product had an amino acid composition shown in Table 2 and an average molecular weight of ca. 900 daltons as shown in Figure 3; no appreciable amounts of free amino acids were present. According to the methionine content (ca. 3%) of this product, it was encoded as OPM₃.

Feeding tests with normal and malnourished rats. By simulating the amino acid composition of OPM₃ (Table 2) we prepared a free amino acid mixture which is encoded as AAM₃. On the other hand, SPI₃ was prepared by mixing free methionine with SPI so as to reach a level of 3% with respect to total methionine content. With these comparable nitrogen sources (SPI₃, OPM₃ and AAM₃) at a nitrogen level of 1.5% each, we formulated three different diets of Harper's composition (18). In a feeding test with normal subjects, male rats (Wister strain) aged 4 weeks, weighing about 80 g each, were fed these experimental diets for 3 weeks. For preparation of protein-malnourished rats as abnormal male rats aged 6 weeks, weighing about 170 g each, were depleted by feeding with a protein-free diet for 2 weeks until their average body weight decreased to approximately 140 g and symptoms of anemia appeared. The malnourished rats were then fed the experimental diets for the following 3 weeks.

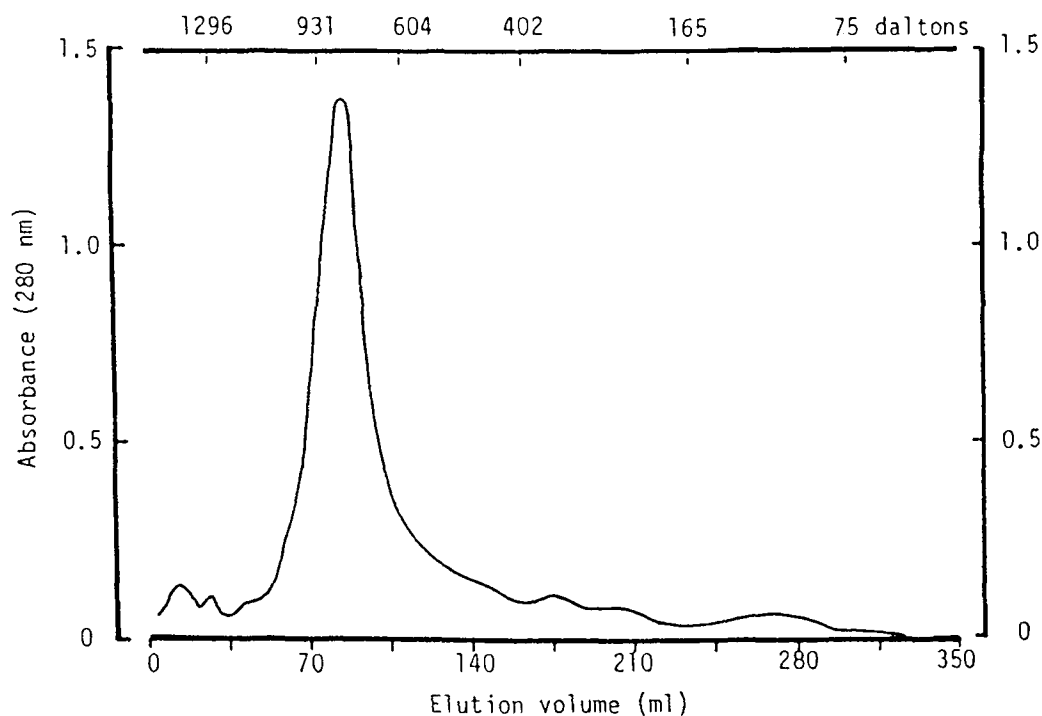


FIG. 3. Chromatography of OPM₃ on Bio-Gel P-2 (2 × 100 cm). The elution was made with 0.2 M phosphate, pH 8, at room temperature. The following markers with known molecular weights (MW) were used: human angiotensin I (MW 1296), human angiotensin III (MW 931), antipain (MW 604), triglutamic acid (MW 402), L-phenylalanine (MW 165), and glycine (MW 75).

RESULTS AND DISCUSSION

Digestion and absorption of EMP₃. It was observed that free amino acids in the portal vein of rats fed the EMP₃ diet reached a maximum at 2 hr after feeding and decreased gradually thereafter. The concentration of free methionine in the portal vein also showed a similar time-course profile. Portal amino acid concentrations of rats fed the simulated AAM diet were almost similar to those observed with the rats of the EMP₃ diet group. These results indicate that the covalently attached methionine occurring in EMP₃ is well bioavailable. However, there was a difference between the EMP diet group and the AAM diet group in terms of the amount of small intestinal nitrogen content. For the EMP₃ diet group we observed a bell-shaped profile showing that the small intestinal nitrogen content reached a maximum at 2 hr after feeding. The amount of total methionine, i.e., free methionine plus peptide-bound methionine, showed a similar time-course profile. However, free methionine was at a very low level over the observation period, representing only 15% of total methionine even at 2 hr after feeding. In particular, the free methionine amount observed for the AAM diet group was six times as high as that observed for the EMP₃ diet group at 1 hr after feeding, whereas in the portal vein this relationship was reversed. Fractionation experiments with the small intestinal content of EMP₃ diet group at 2 hr after feeding showed that it was comprised of molecular weight fractions of higher than 2,000 daltons (18.3%), 1,000–2,000 daltons (21.5%), 500–1,000 daltons (21.1%) and lower than 500 daltons (26.3%). It was also found that a major part of the methionine occurred in the fraction of lower than 500 daltons. Probably, therefore, the methionine existed mostly in the form of oligopeptides.

Supplementation of soy protein with EMP₁₀. The feeding test with rats to examine a supplementary effect of EMP₁₀ gave the following result. The maximum PER value was observed when rats were fed the diet containing the peptide-bound methionine at 0.25–0.30% (Fig. 2). In considering that, in general, the PER value of SPI is maximized when rats are fed SPI whose methionine level is enhanced up to 3–4% by addition of free methionine (20), the result that the methionine level of 2.5–3.0% maximized the PER of SPI (Fig. 2) may

suggest a superior nutritional effect of the peptide-bound methionine occurring in EMP₁₀. Though the most recently published information (21) suggests that soy protein itself needs no supplementary methionine, it may still be useful to apply our method for supplementation to other low-quality proteins for human consumption. It may also be beneficial that EMP₁₀ as well as other EMP products is low in flavor despite its high methionine content.

OPM as a foodstuff for therapeutic use. It is likely that OPM, when administered per os, behaves differently from corresponding AAM in some respects in absorptive and postabsorptive stages (6–7, 22–24), but little is known of any potential difference between them in terms of overall nutritional parameters including body weight gain and PER. For clear detection of such a difference between the nutritional effect of OPM and that of AAM, it may sometimes be preferable to use abnormal animals, e.g., malnourished animals, as subjects for investigation. To obtain information about this point, we carried out a feeding test with normal and protein-malnourished rats where they were fed the OPM₃ diet as a test diet and the SPI₃ and AAM₃ diets as controls. Rats of the different diet groups accepted their diets well. During the feeding period (3 weeks) we measured food intakes and body weight gains to obtain PER values and dietary and fecal nitrogen amounts to obtain digestibilities. We also examined the rats for their blood chemistry involving glutamine-oxoacid transaminase (GOT) and glutamine-pyruvate aminotransferase (GPT) as physiological parameters (25). The results are summarized in Table 3. Both for normal and for malnourished rats, SPI₃ was significantly less digestible than OPM₃. However, both normal and malnourished rats utilized SPI₃ and OPM₃ with similar efficiency. In the case of malnourished rats, the PER value observed for OPM₃ was significantly higher than that observed for AAM₃. It was also likely from the data on GPT that OPM₃ surpassed AAM₃ in blood physiology. These results suggest that OPM₃ could be utilized similarly to or more efficiently than SPI from which it was derived and also that OPM₃ is generally superior to AAM₃. It is thus expected that OPM₃ would be used as an oligopeptide rather than a free amino acid nitrogen source for faster recovery of unhealthy subjects under therapeutic care.

TABLE 3

Performance of a Test with Normal and Protein-Malnourished Rats Fed SPI₃, OPM₃ and AAM₃-Diets for Three Weeks

State of rats	Number of rats	Nitrogen source	Digestibility (%)	Body weight gain (g/day)	PER	GOT (Karmen units)	GPT
Normal	5	SPI ₃	87.0	5.84	2.96	66	8.3
Normal	5	OPM ₃	92.3	6.24	3.12	55	11.0
Normal	5	AAM ₃	92.0	5.81	2.72	69	14.0
Malnourished	5	SPI ₃	87.0	8.10	3.88	68	12.0
Malnourished	5	OPM ₃	91.1	8.67	3.97	55	11.0
Malnourished	5	AAM ₃	92.6	7.82	3.49	65	13.5

^aStudents' t-test. P < 0.05.

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[Received February 19, 1987]