

Hydroxyproline-containing dipeptides and tripeptides quantified at high concentration in human blood after oral administration of gelatin hydrolysate

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Abstract

Several hydroxyproline (Hyp)-containing food-derived collagen peptides were identified in human blood after oral ingestion of gelatin hydrolysates. However, these types of peptides were not quantified in human plasma. In this report, a sensitive LC-MS/MS method was introduced for simultaneous quantitative analysis of Hyp-containing peptides. All peptide concentrations were determined accurately, with all coefficients of determination (r^2) >0.999. The method achieved detection and quantification limits of 0.01 pmol/ml and 12.5–1,000 pmol/ml in plasma, respectively. Concentrations were quantified for nine Hyp-containing peptides in human plasma by this method, identifying Pro-Hyp ($C_{\max} = 60.65 \pm 5.74$ nmol/ml) as the major constituent of food-derived collagen peptides, while the minor components were Ala-Hyp-Gly, Ser-Hyp-Gly, Ala-Hyp, Phe-Hyp, Leu-Hyp, Ile-Hyp, Gly-Pro-Hyp, and Pro-Hyp-Gly (C_{\max} from 23.84 to 0.67 nmol/ml). Thus a total of nine Hyp-containing peptides in human plasma were successfully quantified by this approach. The concentration of Hyp-containing peptides is substantially higher than that following oral administration of other peptides.

Keywords: Collagen, hydroxyproline, plasma, hydroxyproline-containing peptide, quantification

Introduction

Collagen is a major constituent of connective tissues of animals, birds, and fish. Gelatin, a denatured form of collagen, is prepared on an industrial scale from these animals (Shrieber and Seybold 1993). Collagen has a unique triple helix configuration with a repeating sequence (Gly-X-Y)_n, with X and Y being mostly proline and hydroxyproline (Hyp) (Ramshaw and Shah 1998; Bos et al. 1999). Gelatin-based food derivatives obtained from animals, especially fish and pigs, have been attracting worldwide attention as health-food ingredients. Significant amounts of Hyp-containing peptides were found to be present in the peripheral blood of human volunteers after oral ingestion of porcine skin gelatin hydrolysates (Iwai et al. 2005). Recently, some Hyp-containing peptides were also detected in human blood after ingestion of hydrolysate from fish scales (Ohara et al. 2007a). The major constituents of Hyp-containing

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49 peptides that remained in the blood were identified as Ala-Hyp, Pro-Hyp, Ala-Hyp-
50 Gly, Ser-Hyp-Gly, Phe-Hyp, Pro-Hyp-Gly, Gly-Pro-Hyp, Ile-Hyp and Leu-Hyp.
51 These collagen-based peptides represent functional peptides involved in various
52 physiological activities. For example, Pro-Hyp and Gly-Pro-Hyp exert chemotactic
53 effects on fibroblasts, peripheral blood neutrophils (Postlethwaite et al. 1978; Laskin
54 et al. 1986) and monocytes (Postlethwaite and King 1976) in cell culture systems. Gly-
55 Pro-Hyp is also suggested to be involved in platelet aggregation (Knight et al. 1999).
56 Recently, Shigemura et al. (2009) indicated that Pro-Hyp enhanced mice fibroblast cell
57 proliferation. Therefore, it could be assumed that food-derived collagen peptides in
58 blood may be involved in some of the biological activities suggested by animal and
59 human experiments.

60 However, no quantitative analysis of peptides has been reported in an earlier human
61 absorption study. Previous methods to quantify these types of peptides involved
62 subtraction of free Hyp and Hyp-containing peptide concentrations after determining
63 the Hyp concentration in plasma using reverse-phase high-performance liquid chro-
64 matography (Iwai et al. 2005; Aito-Inoue et al. 2006; Ohara et al. 2007a). Thus,
65 quantification of food-derived Hyp-containing peptides has been evaluated by semi-
66 quantitative methods such as determining the recovery of Hyp in each peptide peak.
67 Moreover, it has been difficult to detect and isolate small amounts of food-derived
68 peptides that do not have any marker amino acids or modified amino acids from animal
69 and human blood after oral ingestion.

70 To overcome this problem, a sensitive and convenient liquid chromatography mass
71 spectrometry/mass spectrometry (LC-MS/MS) method was introduced for simulta-
72 neous analysis of Hyp-containing peptides in human plasma after oral ingestion of fish-
73 scale gelatin hydrolysate. Recently, digested mixtures of collagen type II and type I
74 containing many specific peptides and common peptides were analyzed by MS/MS
75 sequencing (Zhang et al. 2006). However, only tetrapeptides to nonapeptides were
76 analyzed to define the collagen type, whereas specific dipeptides and tripeptides from
77 human blood samples were not analyzed.

78 The goal of the present study was to quantify food-derived Hyp-containing peptides
79 in a complex matrix such as human plasma.

82 **Materials and methods**

83 *Gelatin hydrolysate*

84 Enzymatic hydrolysate of fish-scale gelatin was a kind gift from Nitta Gelatin, Ltd
85 (Osaka, Japan). This preparation was of food grade and it can be obtained commer-
86 cially. The average molecular weight of peptides in this gelatin hydrolysate, which did
87 not contain the free form of Hyp, was about 5,000 Da.

89 *Chemicals*

90 Acetonitrile (high-performance liquid chromatography grade), pentafluoropropionic
91 acid, and trichloroacetic acid were purchased from Wako Pure Chemical Industries
92 (Osaka, Japan). Ala-Hyp, Ala-Hyp-Gly, Ser-Hyp-Gly, Pro-Hyp-Gly, Ile-Hyp, Leu-
93 Hyp, and Phe-Hyp were purchased from Kokusan Chemical (Tokyo, Japan), and Pro-
94 Hyp and Gly-Pro-Hyp were purchased from Bachem (Bubendorf, Germany).
95
96

Preparation of standard samples

Standards prepared for nine Hyp-containing peptides (Ala-Hyp, Pro-Hyp, Ala-Hyp-Gly, Ser-Hyp-Gly, Phe-Hyp, Pro-Hyp-Gly, Gly-Pro-Hyp, Ile-Hyp and Leu-Hyp) were dissolved in blank human plasma or water, mixed and diluted to 1 nmol/ml, 5 nmol/ml, 10 nmol/ml, 25 nmol/ml, 50 nmol/ml and 100 nmol/ml. They were then mixed with equal amounts of 5% (w/v) trichloroacetic acid. After filtration with a 4-mm, 0.22- μ m PVDF filter (Millipore, Bedford, MA, USA), 5 μ l of the resulting filtrate was injected into the LC-MS/MS system.

Human study design

The present study was performed according to the Helsinki Declaration and was approved by the Ethical Committee of Meiji Seika Kaisha, Ltd, Food and Health R&D Laboratories. Five healthy male volunteers with no incidence of gelatin allergy (33.0 ± 5.6 years old and 69.8 ± 7.4 kg body weight) participated in the study. Subjects did not consume any food or beverages except for water in the 12-h period prior to the experiment. On the morning of the experiment, the subjects were fasting and each subject orally ingested the fish-scale gelatin hydrolysate concentrate (0.385 g/kg body weight) in water (20% w/v). Three hours after ingestion of the gelatin hydrolysate preparation, the subjects were served a collagen-free lunch, consisting of only a rice ball with salt. Approximately 5 ml venous blood was collected from the cubital vein before (0 h) and 0.5 h, 1 h, 2 h, 4 h, and 7 h after ingesting the hydrolysate. Plasma was obtained after blood centrifugation at 880 $\times g$ for 10 min at 4°C and stored in tubes at -80°C until analysis was performed.

Pre-treatment of blood sample for LC-MS/MS

The plasma was de-proteinized by adding equal amounts of 5% (w/v) trichloroacetic acid. The supernatant was then centrifuged at 14,010 $\times g$ for 10 min at 4°C. After filtering through a 4-mm, 0.22- μ m PVDF filter, 5 μ l of the resulting filtrate was injected into the LC-MS/MS system.

LC-MS/MS analysis

Samples were analyzed by LC-MS/MS. The LC analysis was performed using an ACQUITY UPLC system (Waters, Milford, MA, USA). A particular Octa Decyl Sillica (ODS) column that retains polar compounds tightly was better adapted to this analysis than the conventional ODS column that was used previously. Therefore an ACQUITY UPLC HSS T3 column (2.1 \times 50 mm, 1.7 μ m; Waters) was used for the separation. Gradient elution was carried out with 0.05% (v/v) pentafluoropropionic acid and acetonitrile at a constant flow rate of 0.3 ml/min. The gradient profile with the following proportions (v/v) of acetonitrile was applied (t (min), % acetonitrile): (0 min, 0%), (4 min, 0%), (9 min, 25%), (9.01 min, 80%), (10 min, 80%) (3 min: time was required to reach initial conditions). The column temperature was maintained at 40°C. The Quattro Premier XE tandem quadrupole mass spectrometer was used in positive ion electrospray mode. The ion source was operated at 120°C with a capillary voltage of 3.5 kV. Nitrogen was employed for the desolvation gas at 400°C and 850 l/h. The mode of acquisition was multiple reaction monitoring (MRM) at an argon collision gas pressure of 5.0×10^{-3} mbar. The list of peptides and the MRM transitions, along with

Table I. MRM method parameters.

Peptide	Retention time (min)	MRM transition
Ala-Hyp	1.7	203.3 > 132.1
Pro-Hyp	2.7	229.2 > 70.2
Ala-Hyp-Gly	2.1	260.3 > 189.0
Ser-Hyp-Gly	1.9	276.3 > 189.1
Phe-Hyp	7.8	279.3 > 119.9
Pro-Hyp-Gly	4.0	286.3 > 189.0
Gly-Pro-Hyp	5.6	286.3 > 154.7
Ile-Hyp	7.0	245.3 > 131.9
Leu-Hyp	7.2	245.3 > 131.9

Cone voltage: Pro-Hyp, 25 V; others, 20 V. Collision energy, 15 eV.

255 the retention times, cone voltages, and collision energies for the method, are presented
 256 in Table I. The data were acquired using MassLynx Software version 4.1 (Waters) and
 257 were processed using the TargetLynx application manager.
 258

259 *Pharmacokinetic analysis*

261 Analysis of blood concentration–time data was carried out with a non-compartment
 262 model using WinNonlin Professional (version 5.2.1; Pharsight Co., Mountain
 263 View, CA, USA). The total area under the concentration–time curve ($AUC_{0-7\text{ h}}$)
 264 was calculated by the trapezoidal rule based on the plasma concentrations up to the
 265 time of final measurement using the WinNonlin Professional program.
 266

267 **Results**

269 *Analysis of standards*

270 Figure 1 shows typical MRM chromatograms of the nine Hyp-containing peptide
 271 standards. The total run-time per sample was only 13 min. The sensitivity of the method
 272 was evaluated by determining the limit of detection (LOD) and the limit of quantification
 273 (LOQ). The LOD was defined as the concentration of the nine Hyp-containing peptides
 274 with a signal-to-noise ratio of 3, for the chromatographic peaks from 0.01 pmol/ml to
 275 100 nmol/ml, stepwise. The LOQ was the lowest standard concentration with a signal-to-
 276 noise ratio of 10. The LOD and LOQ for a 5 μ l injection, coefficients of determination and
 277 recovery for each of the nine Hyp-containing peptides in plasma are presented in Table II.
 278 The method achieved detection and quantification limits of 0.01 pmol/ml and 12.5–1,000
 279 pmol/ml in plasma, respectively. The LOQ was as follows: Ala-Hyp, 225 pmol/ml;
 280 Ser-Hyp-Gly, 125 pmol/ml; Ala-Hyp-Gly, 200 pmol/ml; Pro-Hyp, 1000 pmol/ml;
 281 Pro-Hyp-Gly, 125 pmol/ml; Gly-Pro-Hyp, 75 pmol/ml; Ile-Hyp, 50 pmol/ml; Leu-Hyp,
 282 12.5 nmol/ml; and Phe-Hyp; 150 pmol/ml.
 283

284 The linearity of the method was investigated by spiking blank human plasma (obtained
 285 before collagen ingestion) with known concentrations of the nine Hyp-containing
 286 peptides at six concentration levels ranging from 1 to 100 nmol/ml. The linearity of
 287 measurement over the calibration curve range was good for all peptides measured, and all
 288 coefficients of determination (r^2) were >0.999. Furthermore, the recovery of standards
 289 added to blank human plasma (obtained before collagen ingestion) was investigated with
 25 nmol/ml of the nine Hyp-containing peptides, and their recovery rates were 97–100%

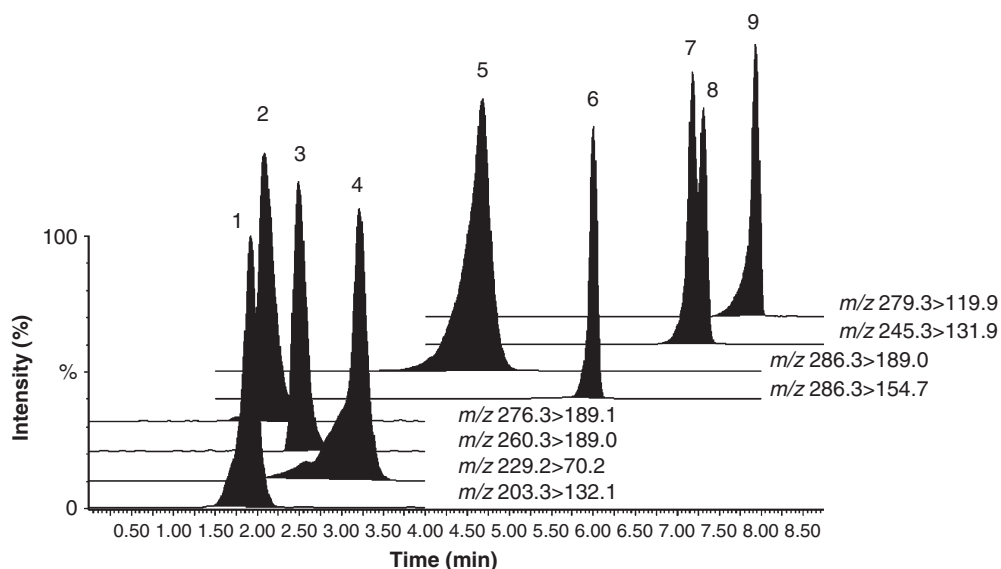


Figure 1. MRM chromatogram of nine Hyp-containing peptides. Peak 1, Ala-Hyp (m/z 203.3 > 132.1); peak 2, Ser-Hyp-Gly (m/z 276.3 > 189.1); peak 3, Ala-Hyp-Gly (m/z 260.3 > 189.0); peak 4, Pro-Hyp (m/z 229.2 > 70.2); peak 5, Pro-Hyp-Gly (m/z 286.3 > 189.0); peak 6, Gly-Pro-Hyp (m/z 286.3 > 154.7); peak 7, Ile-Hyp (m/z 245.3 > 131.9); peak 8, Leu-Hyp (m/z 245.3 > 131.9); peak 9, Phe-Hyp (m/z 279.3 > 119.9).

290 (Table II). In addition, other concentrations of the nine Hyp-containing peptides at
 291 1 nmol/ml, 5 nmol/ml, 10 nmol/ml, 50 nmol/ml and 100 nmol/ml were investigated.
 292 Their recovery rates were 94–107% (data not shown). Therefore, this method is
 293 adequate to detect these nine Hyp-containing peptides.

294 295 296 *Levels of nine Hyp-containing peptides in human plasma*

297 Figure 2 shows the amounts of the nine Hyp-containing peptides in human plasma after
 298 oral ingestion of fish-scale gelatin hydrolysate. Only negligible amounts of each peptide
 299 were observed before the ingestion of fish-scale gelatin hydrolysate. In all subjects, the nine
 300 Hyp-containing peptides in the plasma increased after oral ingestion and reached a

Table II. Correlation coefficient, recovery, limit of quantification, and detection data obtained from LC-MS/MS analysis of nine Hyp-containing peptides in human plasma ($n = 6$).

Peptide	Correlation coefficient	Percentage recovery (% relative standard deviation)	LOQ (pmol/ml)	LOD (pmol/ml)
Ala-Hyp	0.999	100 (2)	225	0.01
Ser-Hyp-Gly	0.999	99 (1)	125	0.01
Ala-Hyp-Gly	0.999	99 (3)	200	0.01
Pro-Hyp	0.999	100 (5)	1,000	0.01
Pro-Hyp-Gly	0.999	98 (4)	125	0.01
Gly-Pro-Hyp	0.999	99 (2)	75	0.01
Ile-Hyp	0.999	97 (1)	50	0.01
Leu-Hyp	0.999	99 (1)	12.5	0.01
Phe-Hyp	0.999	99 (3)	150	0.01

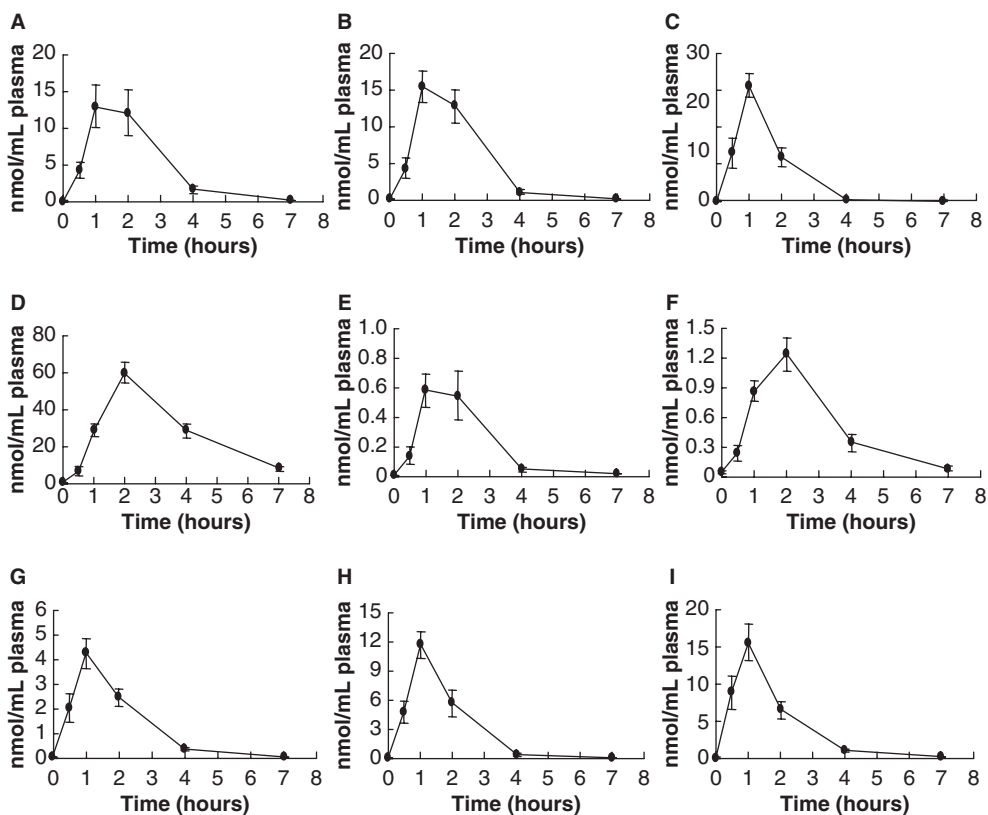


Figure 2. Plasma levels of nine Hyp-containing peptides after oral ingestion of fish-scale gelatin hydrolysate. (a) Ala-Hyp; (b) Ser-Hyp-Gly; (c) Ala-Hyp-Gly; (d) Pro-Hyp; (e) Pro-Hyp-Gly; (f) Gly-Pro-Hyp; (g) Ile-Hyp; (h) Leu-Hyp; (i) Phe-Hyp. Values presented as the mean \pm standard error, $n = 5$ subjects.

301 maximum 1–2 h after ingestion. The T_{\max} (h), C_{\max} (nmol/ml), and AUC (h nmol/ml) of
 302 the nine Hyp-containing peptides are presented in Table III. The T_{\max} values for Pro-Hyp
 303 and Gly-Pro-Hyp were reached 2 h after oral ingestion of fish-scale gelatin hydrolysate. On
 304 the other hand, the T_{\max} values for the seventh through ninth Hyp-containing peptides
 305 were from 1 to 1.6 h after oral ingestion of the hydrolysate. The C_{\max} in plasma was
 306 60.65 ± 5.74 nmol/ml plasma, and the C_{\max} of Pro-Hyp was higher than that of the other
 307 eight Hyp-containing peptides. The calculated AUC_{0-7h} of each Hyp-containing peptide
 308 was as follows: Ala-Hyp, 34.55 ± 8.48 h nmol/ml; Ser-Hyp-Gly, 36.25 ± 5.26 h nmol/ml;
 309 Ala-Hyp-Gly, 37.72 ± 3.98 h nmol/ml; Pro-Hyp, 201.17 ± 18.78 h nmol/ml; Pro-Hyp-Gly,
 310 1.49 ± 0.31 h nmol/ml; Gly-Pro-Hyp, 3.62 ± 0.57 h nmol/ml; Ile-Hyp, 9.06 ± 1.19 h nmol/
 311 ml; Leu-Hyp, 21.30 ± 3.36 h nmol/ml; and Phe-Hyp, 28.85 ± 4.50 h nmol/ml. This result
 312 indicated that Pro-Hyp was the major Hyp-containing peptide in plasma after oral
 313 ingestion of fish-scale gelatin hydrolysate, as reported earlier (Ohara et al. 2007a).

314 Discussion

315 Several Hyp-containing food-derived collagen peptides were identified in human blood
 316 after oral ingestion of gelatin hydrolysates. However, none of these peptides were
 317

Table III. AUC_{0-7 h} of nine Hyp-containing peptides in human plasma after oral ingestion of fish-scale gelatin hydrolysate.

Peptide	T _{max} (h)	C _{max} (nmol/ml)	AUC _{0-7 h}
Ala-Hyp	1.60 ± 0.24	13.70 ± 2.78	34.55 ± 8.48
Ser-Hyp-Gly	1.40 ± 0.24	16.58 ± 1.72	36.25 ± 5.26
Ala-Hyp-Gly	1.00 ± 0.00	23.84 ± 2.44	37.72 ± 3.98
Pro-Hyp	2.00 ± 0.00	60.65 ± 5.74	201.17 ± 18.78
Pro-Hyp-Gly	1.40 ± 0.24	0.67 ± 0.14	1.49 ± 0.31
Gly-Pro-Hyp	2.00 ± 0.00	1.24 ± 0.17	3.62 ± 0.57
Ile-Hyp	1.00 ± 0.00	4.26 ± 0.60	9.06 ± 1.19
Leu-Hyp	1.00 ± 0.00	11.71 ± 1.35	21.30 ± 3.36
Phe-Hyp	1.00 ± 0.00	15.61 ± 2.46	28.85 ± 4.50

Values presented as the mean ± standard error, *n* = 5 subjects.

318 quantified in human plasma. In this report, a LC-MS/MS method was introduced to
 319 quantify Hyp-containing peptides in human plasma after oral ingestion of fish-scale
 320 gelatin hydrolysate. The recovery of standards added to plasma was quantified,
 321 confirming that this method could be used to measure concentrations of Hyp-contain-
 322 ing peptides without derivatization. In addition, the linearity of the measurements was
 323 evaluated, and results confirmed that it was accurate over the calibration curve range
 324 for all peptides. Previous approaches to measuring peptides containing Hyp were based
 325 on their derivatization with phenyl isothiocyanate (Iwai et al. 2005; Ohara et al.
 326 2007a; Aito-Inoue et al. 2006).

327 The major constituent of food-derived collagen peptides remaining in blood was
 328 confirmed to be Pro-Hyp (AUC_{0-7 h} = 201.17 ± 18.78 h nmol/ml), while the minor
 329 components were Ala-Hyp-Gly, Ser-Hyp-Gly, Ala-Hyp, Phe-Hyp, Leu-Hyp, Ile-Hyp,
 330 Gly-Pro-Hyp, and Pro-Hyp-Gly (AUC_{0-7 h} from 37.72 to 1.49 h nmol/ml). This result
 331 indicated that Pro-Hyp was the major Hyp-containing peptide in plasma after oral
 332 ingestion of fish-scale gelatin hydrolysate, as reported earlier (Ohara et al. 2007a). In
 333 the present study, Pro-Hyp reached its maximum concentration in plasma 2 h after oral
 334 ingestion of fish-scale gelatin hydrolysate, while Ala-Hyp and Ala-Hyp-Gly reached their
 335 maximum concentrations 1 h after ingestion of the hydrolysate. Another study reported
 336 that more than 75% of Pro-Hyp remained 24 h after being added *in vitro* to human serum
 337 (Iwai et al. 2005). Therefore, Pro-Hyp can be considered indigestible by human blood.

338 It is well known that the abundance of the oligopeptide transporter (PEPT-1) in the
 339 brush-border membrane of the intestinal epithelium is the principal mechanism for
 340 regulation of transport of products of protein digestion (dipeptides and tripeptides).
 341 Gly-Pro-Hyp can be partially hydrolyzed by the brush-border membrane-bound
 342 aminopeptidase N to remove Gly, and the resulting Pro-Hyp may be transported
 343 into small intestinal epithelial cells via the H⁺-coupled PEPT-1 (Aito-Inoue et al.
 344 2007). It therefore may be possible for Hyp-containing dipeptides or tripeptides to be
 345 absorbed transcellularly, at least partly, via this peptide transporter (Adibi 2003).

346 After peptide ingestion, dipeptides were detected in human blood, but their con-
 347 centrations were quite low. Matsui et al. (2002) reported that the dipeptide Val-Tyr was
 348 observed in plasma 2 h after oral peptide administration. The maximal Val-Tyr
 349 concentration in plasma was 2,041 ± 148 fmol/ml. Morifuji et al. (2009) reported
 350 the plasma levels of Val-Leu, Ile-Leu and Leu-Leu after ingestion of soy and whey
 351 protein hydrolysate. The maximal Val-Leu, Ile-Leu and Leu-Leu concentrations in

407 plasma were 25 nmol/l, 40 nmol/l and 6 nmol/l, respectively. In the present study, Pro-
408 Hyp was the major Hyp-containing peptide in plasma after oral ingestion of fish-scale
409 gelatin hydrolysate, and the maximal level in plasma was 60.65 ± 5.74 nmol/ml plasma.
410 The C_{\max} of Pro-Hyp was higher than that of Val-Tyr. Stimulation of human fibroblast
411 proliferation and hyaluronan synthesis by Pro-Hyp has been achieved at a concentra-
412 tion of 100 nmol/ml (Ohara et al. 2007b). The amount of Pro-Hyp in plasma 2 h after
413 oral ingestion of fish-scale gelatin hydrolysate is approximately 60 nmol/ml plasma.
414 Therefore, the total Pro-Hyp content in plasma or skin is estimated to reach approx-
415 imately 100 nmol/ml. This suggests that oral ingestion of collagen can result in
416 biological activities that depend on food-derived Hyp-containing peptides.
417

418 **Conclusions**

419 Concentrations of nine Hyp-containing peptides were determined in human plasma
420 after oral ingestion of fish-scale gelatin hydrolysate. Pro-Hyp was the major constituent
421 of food-derived collagen peptides, while the minor components were Ala-Hyp-Gly,
422 Ser-Hyp-Gly, Ala-Hyp, Phe-Hyp, Leu-Hyp, Ile-Hyp, Gly-Pro-Hyp, and Pro-Hyp-Gly.
423 The concentration of Hyp-containing peptides is substantially higher than that
424 following oral ingestion of other peptides.
425

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