

NH₂-Terminally Modified Gastric Inhibitory Polypeptide Exhibits Amino-Peptidase Resistance and Enhanced Antihyperglycemic Activity

Finbarr P.M. O'Harte, Mark H. Mooney, and Peter R. Flatt

Gastric inhibitory polypeptide (GIP) is an important insulin-releasing hormone of the enteroinsular axis that, like glucagon-like peptide 1(7-36) amide (tGLP-1), has a functional profile of possible therapeutic value for type 2 diabetes. Both incretin hormones are rapidly inactivated in plasma by the exopeptidase dipeptidyl peptidase (DPP) IV. The present study examined the ability of NH₂-terminal modification of human GIP to protect from plasma degradation and enhance insulin-releasing and antihyperglycemic activity. Degradation of GIP by incubation at 37°C with purified DPP IV was clearly evident after 4 h (54% intact). After 12 h, >60% of GIP was converted to GIP(3-42), whereas >99% of NH₂-terminally modified Tyr¹-glucitol GIP remained intact. Tyr¹-glucitol GIP was similarly resistant to serum degradation. The formation of GIP(3-42) was almost completely abolished by inhibition of plasma DPP IV with diprotin A. Effects of GIP and Tyr¹-glucitol GIP were examined in Wistar rats after intraperitoneal injection of either peptide (10 nmol/kg) together with glucose (18 mmol/kg). Plasma glucose concentrations were significantly lower and insulin concentrations higher after both peptides compared with glucose alone. More importantly, individual glucose values at 15 and 30 min together with the areas under the curve (AUCs) for glucose were significantly lower after administration of Tyr¹-glucitol GIP compared with GIP (AUC 255 ± 33 vs. 368 ± 8 mmol · l⁻¹ · min⁻¹, respectively; *P* < 0.01). This was associated with a significantly greater and more protracted insulin response after Tyr¹-glucitol GIP than GIP (AUC 773 ± 41 vs. 639 ± 39 ng · ml⁻¹ · min⁻¹; *P* < 0.05). These data demonstrate that Tyr¹-glucitol GIP displays resistance to plasma DPP IV degradation and exhibits enhanced antihyperglycemic activity and insulin-releasing action in vivo. *Diabetes* 48:758-765, 1999

From the School of Biomedical Sciences, University of Ulster, Cromore Road, Coleraine, Northern Ireland.

Address correspondence and reprint requests to Dr. Finbarr O'Harte, School of Biomedical Sciences, University of Ulster, Coleraine, N. Ireland, BT52 1SA. E-mail: fpm.oharte@ulst.ac.uk.

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AUC, area under the curve; DPA, diprotin A; DPP, dipeptidyl peptidase; ESI-MS, electrospray ionization mass spectrometry; GIP, gastric inhibitory polypeptide; GLP-1, glucagon-like peptide 1; HPLC, high-performance liquid chromatography; RIA, radioimmunoassay; TFA, trifluoroacetic acid; tGLP-1, glucagon-like peptide-1(7-36) amide.

Gastric inhibitory polypeptide (GIP) and glucagon-like peptide-1(7-36) amide (tGLP-1) are two important insulin-releasing hormones secreted from endocrine cells in the intestinal tract in response to feeding (1,2). Together with autonomic nerves, they play a vital role supporting the pancreatic islets in the control of blood glucose homeostasis and nutrient metabolism (1,3).

Dipeptidyl peptidase (DPP) IV (EC 3.4.14.5) has been identified as a key enzyme responsible for inactivation of GIP and tGLP-1 in serum (4,5). DPP IV is completely inhibited in serum by the addition of diprotin A (DPA) (0.1 mmol/l) (4). This inactivation occurs through the rapid removal of the NH₂-terminal dipeptides Tyr¹-Ala² and His⁷-Ala⁸, giving rise to the main metabolites GIP(3-42) and glucagon-like peptide 1(9-36) amide, respectively. These truncated peptides are reported to lack biological activity or even to serve as antagonists at GIP or tGLP-1 receptors (6-9). The resulting biological half-lives of these incretin hormones in vivo are therefore very short, estimated to be no longer than ~5 min (5,10-12). In situations of normal glucose regulation and pancreatic β-cell sensitivity, this short duration of action is advantageous in facilitating momentary adjustments to homeostatic control. However, the current goal of a possible therapeutic role of incretin hormones, particularly tGLP-1, in type 2 diabetes therapy is frustrated by a number of factors in addition to finding a convenient route of administration (13). Most notable of these are rapid peptide degradation and rapid absorption (peak concentrations reached at 20 min) and the resulting need for both high dosage and precise timing with meals (13-15). Recent therapeutic strategies have focused on precipitated preparations to delay peptide absorption (16) and inhibition of glucagon-like peptide 1 degradation using specific inhibitors of DPP IV (17-19). A possible therapeutic role is also suggested by the observation that a specific inhibitor of DPP IV, isoleucine thiazolidide, lowered blood glucose and enhanced insulin secretion in glucose-treated diabetic obese Zucker rats, presumably by protecting against catabolism of the incretin hormones tGLP-1 and GIP (18).

Numerous studies have indicated that tGLP-1 infusion restores pancreatic β-cell sensitivity, insulin secretory oscillations, and improved glycemic control in various groups of patients with impaired glucose tolerance (IGT) or type 2 diabetes (13,15,20-22). Longer-term studies also show significant benefits of tGLP-1 injections in type 2 and possibly type 1 diabetes therapy (20,23,24), providing a major incentive to develop an orally effective or long-acting tGLP-1 analog (13).

Several attempts have been made to produce structurally modified analogs of tGLP-1 that are resistant to DPP IV degradation (25–27). A significant extension of serum half-life is observed with His⁷-glucitol tGLP-1 and tGLP-1 analogs substituted at position 8 with Gly, Aib, Ser, or Thr (25–27). These structural modifications appear to impair receptor binding and insulinotropic activity, however, thereby compromising part of the benefit of protection from proteolytic degradation (25–28). Thus, in our own recent studies using His⁷-glucitol tGLP-1, resistance to DPP IV and serum degradation was accompanied by severe loss of insulin-releasing activity (26,28).

GIP shares not only the same degradation pathway as tGLP-1 but many similar physiologic actions as well, including stimulation of insulin and somatostatin secretion and enhancement of glucose disposal (1). These actions are viewed as key aspects in the antihyperglycemic properties of tGLP-1 (13), and there is therefore good expectation that GIP may have similar potential in type 2 diabetes therapy. Indeed, compensation by GIP is held to explain the modest disturbances of glucose homeostasis observed in tGLP-1 knockout mice (29). Apart from early studies (30), the antidiabetic potential of GIP has not been explored, and tGLP-1 may seem more attractive, since it is viewed by some as a more potent insulin secretagogue when infused at so-called physiologic concentrations estimated by radioimmunoassay (RIA) (31).

In a recent study, we have shown that NH₂-terminal glycation of GIP markedly enhances the insulin-releasing effect of the peptide on clonal β-cells (32). If such structural modification also confers DPP IV resistance, the potential attractiveness of this peptide for possible type 2 diabetes therapy would be considerably enhanced. The present study has explored this issue by examining *in vitro* degradation of Tyr¹-glucitol GIP together with evaluation of its antihyperglycemic and insulin-releasing properties *in vivo*. The results demonstrate clearly that this novel GIP analog exhibits a substantial resistance to aminopeptidase degradation and increased glucose-lowering activity compared with the native human GIP.

RESEARCH DESIGN AND METHODS

Materials. Human GIP was purchased from American Peptide Company (Sunnyvale, CA). High-performance liquid chromatography (HPLC)-grade acetonitrile was obtained from Rathburn (Walkersburn, Scotland). Sequencing grade trifluoroacetic acid (TFA) was obtained from Aldrich (Poole, Dorset, U.K.). All other chemicals purchased, including Dextran T-70, activated charcoal, sodium cyanoborohydride, and bovine serum albumin fraction V, were from Sigma (Poole, Dorset, U.K.). DPA was purchased from Calbiochem-Novabiochem (Beeston, Nottingham, U.K.), and rat insulin standard for RIA was obtained from Novo Industri (Copenhagen, Denmark). Reverse-phase Sep-Pak cartridges (C-18) were purchased from Millipore (Milford, MA). All water used in these experiments was purified using a Milli-Q Water Purification System (Millipore, Milford, MA).

Preparation of Tyr¹-glucitol GIP. Tyr¹-glucitol GIP was prepared and purified by HPLC as described previously (32). In brief, human GIP was incubated with glucose under reducing conditions in 10 mmol/l sodium phosphate buffer at pH 7.4 for 24 h. The reaction was stopped by addition of 0.5 mol/l acetic acid (30 μl), the mixture was applied to a Vydac (C-18) (4.6 × 250 mm) analytical HPLC column (The Separations Group, Hesperia, CA), and gradient elution conditions were established using aqueous/TFA and acetonitrile/TFA solvents, as described previously. Fractions corresponding to the glycosylated peaks were pooled, taken to dryness under vacuum using an AES 1000 Speed-Vac concentrator (Life Sciences International, Runcorn, U.K.), and purified to homogeneity on a Supelcosil (C-8) (4.6 × 150 mm) column (Supelco, Poole, Dorset, U.K.).

Degradation of GIP and Tyr¹-glucitol GIP by DPP IV. HPLC-purified GIP or Tyr¹-glucitol GIP were incubated at 37°C with DPP-IV (5 mU) for various time periods in a reaction mixture made up to 500 μl with 50 mmol/l triethanolamine-HCl, pH 7.8 (final peptide concentration 1 μmol/l) (4). Enzymatic reactions were terminated after 0, 2, 4, and 12 h by addition of 5 μl of 10% (vol/vol) TFA/water. Sam-

ples were made up to a final volume of 1.0 ml with 0.12% (vol/vol) TFA and stored at –20°C before HPLC analysis.

Degradation of GIP and Tyr¹-glucitol GIP by human plasma. Pooled human plasma (20 μl) taken from six healthy fasted male subjects was incubated at 37°C with GIP or Tyr¹-glucitol GIP (10 μg) for 0 and 4 h in a reaction mixture made up to 500 μl, containing 50 mmol/l triethanolamine/HCl buffer, pH 7.8. Incubations for 4 h were also performed in the presence of DPA (5 mU). The reactions were terminated by addition of 5 μl TFA, and the final volume was adjusted to 1.0 ml with 0.1% (vol/vol) TFA/water. Samples were centrifuged (13,000g, 5 min), and the supernatant was applied to a C-18 Sep-Pak cartridge (Millipore) that was previously primed and washed with 0.1% (vol/vol) TFA/water. After washing with 20 ml 0.12% TFA/water, bound material was released by elution with 2 ml of 80% (vol/vol) acetonitrile/water and concentrated using a Speed-Vac concentrator (Life Sciences International). The volume was adjusted to 1.0 ml with 0.12% (vol/vol) TFA/water before HPLC purification.

HPLC analysis of degraded GIP and Tyr¹-glucitol GIP. Samples were applied to a Vydac C-18 wide-pore column equilibrated with 0.12% (vol/vol) TFA/water at a flow rate of 1.0 ml/min. Using 0.1% (vol/vol) TFA in 70% acetonitrile/water, the concentration of acetonitrile in the eluting solvent was raised from 0 to 31.5% over 15 min, to 38.5% over 30 min, and to 70% over 5 min using linear gradients. The absorbance was monitored at 206 nm, and peak areas were evaluated using a model 2221 LKB integrator. Samples recovered manually were concentrated using a Speed-Vac concentrator.

Electrospray ionization mass spectrometry. Samples for electrospray ionization mass spectrometry (ESI-MS) analysis containing intact and degradation fragments of GIP (from DPP IV and plasma incubations) as well as Tyr¹-glucitol GIP were further purified by HPLC. Peptides were dissolved (~400 pmol) in 100 μl water and applied to the LCQ benchtop mass spectrometer (Finnigan MAT, Hemel Hempstead, U.K.) equipped with a microbore C-18 HPLC column (150 × 2.0 mm, Phenomenex, Macclesfield, U.K.). Samples (30 μl direct-loop injection) were injected at a flow rate of 0.2 ml/min under isocratic conditions in 35% (vol/vol) acetonitrile/water. Mass spectra were obtained from the quadrupole ion trap mass analyzer and recorded. Spectra were collected using full-ion scan mode over the mass-to-charge (*m/z*) range 150–2,000. The molecular masses of GIP and related structures were determined from ESI-MS profiles using prominent multiple charged ions and the equation $M_r = iM_i - iM_H$, where M_r is molecular mass; M_i is *m/z* ratio; i is number of charges; and M_H is mass of a proton.

***In vivo* biological activity of GIP and Tyr¹-glucitol GIP.** Effects of GIP and Tyr¹-glucitol GIP on plasma glucose and insulin concentrations were examined using 10- to 12-week-old male Wistar rats. The animals were housed individually in an air-conditioned room at 22 ± 2°C with a 12-h light/12-h dark cycle. Drinking water and a standard rodent maintenance diet (Trouw Nutrition, Belfast, Northern Ireland) were supplied *ad libitum*. Food was withdrawn for an 18-h period before intraperitoneal injection of glucose (18 mmol/kg body wt), alone or in combination with either GIP or Tyr¹-glucitol GIP (10 nmol/kg). Test solutions were administered in a final volume of 8 ml/kg body wt. Blood samples were collected at 0, 15, 30, and 60 min from the cut tip of the tail of conscious rats into chilled fluoride/heparin microcentrifuge tubes (Sarstedt, Nümbrecht, Germany). Samples were centrifuged using a Beckman microcentrifuge for 30 s at 13,000g. Plasma samples were divided into aliquots and stored at –20°C before glucose and insulin determinations. All animal studies were done in accordance with the Animals (Scientific Procedures) Act 1986.

Analyses. Plasma glucose was assayed by an automated glucose oxidase procedure using a Beckman Glucose Analyzer II (33). Plasma insulin was determined by dextran charcoal radioimmunoassay as described previously (34). Incremental areas under plasma glucose and insulin curves (AUCs) were calculated using a computer program (CAREA) employing the trapezoidal rule (35) with baseline subtraction. Results are expressed as mean ± SE, and values were compared using the Student's unpaired *t* test. Groups of data were considered to be significantly different if *P* < 0.05.

RESULTS

Degradation of GIP and Tyr¹-glucitol GIP by DPP IV. Figure 1 illustrates the typical peak profiles obtained from the HPLC separation of the products obtained from the incubation of GIP (left panels) or Tyr¹-glucitol GIP (right panels) with DPP IV for 0, 2, 4, and 12 h. The retention times of GIP and Tyr¹-glucitol GIP at time 0 were 21.93 and 21.75 min, respectively. Degradation of GIP was evident after 4-h incubation (54% intact), and by 12 h the majority (60%) of intact GIP was converted to the single product, with a retention time of 21.61 min. Tyr¹-glucitol GIP remained almost completely intact throughout 2- to 12-h incubation.

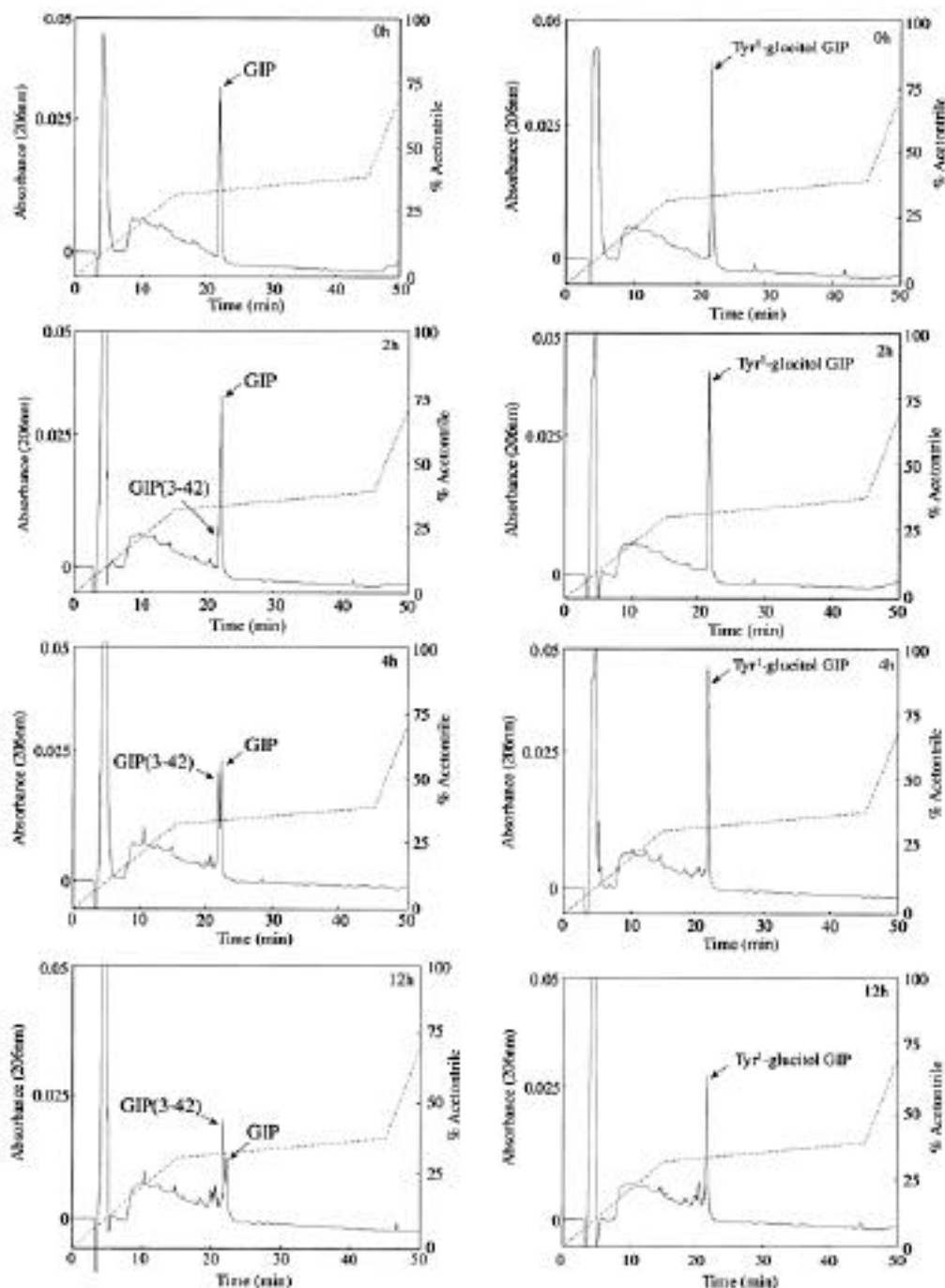


FIG. 1. Degradation of GIP and Tyr¹-glucitol GIP by DPP IV. Representative HPLC profiles obtained after incubation of GIP (*left panels*) or Tyr¹-glucitol GIP (*right panels*) with DPP IV for 0, 2, 4, and 12 h. Incubations of GIP and Tyr¹-glucitol GIP exposed to DPP IV were separated on a Vydac C-18 column using linear gradients 0–31.5% acetonitrile over 15 min, to 38.5% over 30 min, and to 70% over 5 min. Left panels show HPLC profiles of intact GIP (retention time 21.93 min) and GIP(3-42) (retention time 21.61 min). Right panels show HPLC profiles obtained for Tyr¹-glucitol GIP (retention time 21.75 min). HPLC peaks corresponding to intact GIP, GIP(3-42), and Tyr¹-glucitol GIP are indicated.

Degradation of GIP and Tyr¹-glucitol GIP by human plasma. Figure 2 shows a set of typical HPLC profiles of the products obtained from the incubation of GIP or Tyr¹-glucitol GIP with human plasma for 0 and 4 h GIP (left panels). With a retention time of 22.06 min, GIP was readily metabolized by plasma within 4-h incubation, giving rise to the appearance of a major degradation peak with a retention time of 21.74 min. In contrast, the incubation of Tyr¹-glucitol GIP under similar conditions (right panels) did not result in

the formation of any detectable degradation fragments during this time, with only a single peak being observed at retention time 21.77 min. Addition of DPA, a specific inhibitor of DPP IV, to GIP during the 4-h incubation completely inhibited degradation of the peptide by plasma.

Identification of GIP degradation fragments by ESI-MS. Figure 3 shows the monoisotopic molecular masses obtained for GIP (*A*), Tyr¹-glucitol GIP (*B*), and the major plasma degradation fragment of GIP (*C*) using ESI-MS. The peptides

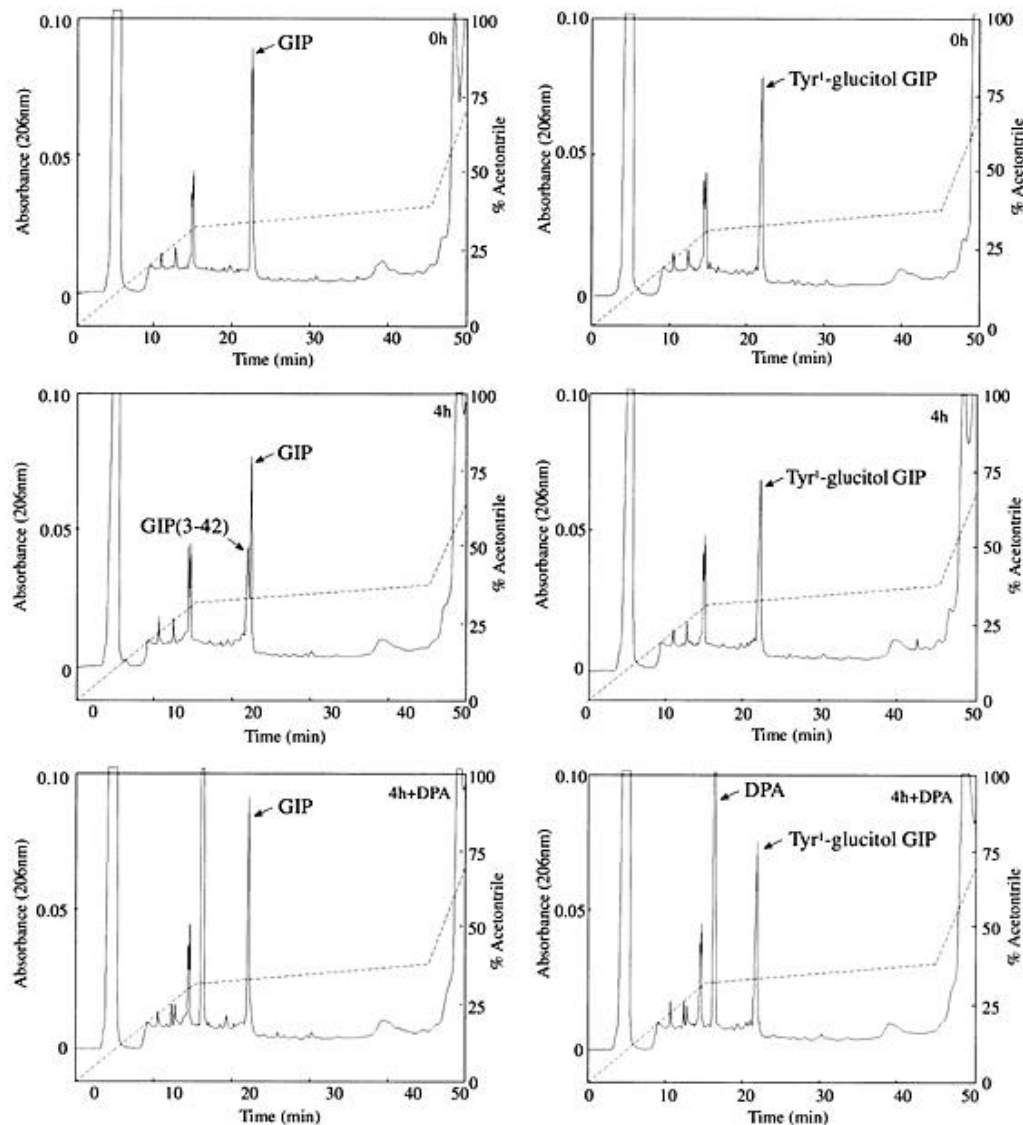


FIG. 2. Degradation of GIP and Tyr¹-glucitol GIP by human plasma. Representative HPLC profiles obtained after incubation of GIP (*left panels*) and Tyr¹-glucitol GIP (*right panels*) with human plasma for 0 and 4 h and for 4 h in the presence of 5 μM of DPA. GIP and Tyr¹-glucitol GIP incubations were separated with a Vydac C-18 column using linear gradients 0–31.5% acetonitrile over 15 min, to 38.5% over 30 min and to 70% over 5 min. Peaks corresponding to intact GIP, GIP(3-42), and Tyr¹-glucitol GIP are indicated. A major peak corresponding to the specific DPP IV inhibitor tripeptide DPA appears in the bottom panels, with retention time 16.29 min.

analyzed were purified from plasma incubations as shown in Fig. 2. The exact molecular mass (M_r) of the peptides were calculated using the equation $M_r = iM_i - iM_h$ as defined in METHODS. After spectral averaging was performed, prominent multiple charged species ($M+3H$)³⁺ and ($M+4H$)⁴⁺ were detected from GIP at m/z 1661.6 and 1246.8, corresponding to intact M_r 4981.8 and 4983.2 Da, respectively (Fig. 3A). Similarly for Tyr¹-glucitol GIP, ($M+4H$)⁴⁺ and ($M+5H$)⁵⁺ were detected at m/z 1287.7 and 1030.3, corresponding to intact molecular masses of M_r 5146.8 and 5146.5 Da, respectively (Fig. 3B). The difference between the observed molecular masses of the quadruply charged GIP and the NH₂-terminally modified GIP species (163.6 Da) indicated that the latter peptide contained a single glucitol adduct corresponding to Tyr¹-glucitol GIP. Figure 3C shows the prominent multiply charged species ($M+3H$)³⁺ and ($M+4H$)⁴⁺ detected from the major fragment of GIP at m/z 1583.8 and 1188.1, corresponding to intact

M_r 4748.4 and 4748.4 Da, respectively. This corresponds with the theoretical mass of the NH₂-terminally truncated form of the peptide GIP(3-42). This fragment was also the major degradation product of DPP IV incubations (data not shown).

Effects of GIP and Tyr¹-glucitol GIP on plasma glucose homeostasis. Figures 4 and 5 show the effects of intraperitoneal glucose alone (18 mmol/kg) (control group) and glucose in combination with GIP or Tyr¹-glucitol GIP (10 nmol/kg) on plasma glucose and insulin concentrations. Compared with the control group, plasma glucose concentrations and AUCs were significantly lower after administration of either GIP or Tyr¹-glucitol GIP (Fig. 4A and B). Furthermore, individual values at 15 and 30 min together with AUCs were significantly lower after administration of Tyr¹-glucitol GIP compared with GIP. Consistent with the established insulin-releasing properties of GIP, plasma insulin concentrations of both peptide-treated groups were significantly raised at 15 and 30 min com-

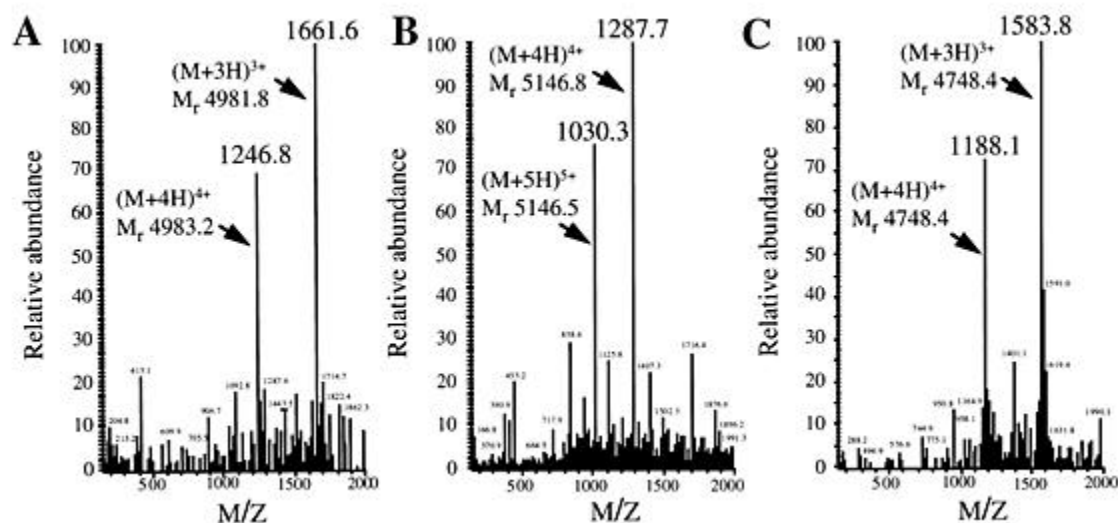


FIG. 3. ESI-MS of GIP, Tyr¹-glucitol GIP, and the major degradation fragment GIP(3-42). Samples containing GIP (A), Tyr¹-glucitol GIP (B), and the major degradation fragment of GIP [GIP(3-42)] (C) isolated from plasma incubations (Fig. 2). Peptides were dissolved (~400 pmol) in 100 μ l water and applied to the LC/MS equipped with a microbore C-18 HPLC column. Samples (30- μ l direct-loop injection) were applied at a flow rate of 0.2 ml/min under isocratic conditions (35% acetonitrile/water). Mass spectra were recorded using a quadrupole ion trap mass analyzer. Spectra were collected using full-ion scan mode over the m/z range 150–2,000. The M_r of GIP and related structures were determined from ESI-MS profiles using prominent multiple charged ions and the following equation: $M_r = iM_i - iM_h$ (see METHODS).

pared with the values after administration of glucose alone (Fig. 5A). The overall insulin responses, estimated as AUCs, were also significantly greater for the two peptide-treated groups (Fig. 5B). Despite lower prevailing glucose concentrations than the GIP-treated group, plasma insulin response, calculated as AUC, after Tyr¹-glucitol GIP was significantly greater than after GIP (Fig. 5B). The significant elevation of plasma insulin at 30 min is of particular interest, suggesting that the insulin-releasing action of Tyr¹-glucitol GIP is more pro-

tracted than GIP even in the face of a diminished glycemic stimulus (Figs. 4A and 5A).

DISCUSSION

The 42-amino acid GIP is an important incretin hormone released into the circulation from endocrine K-cells of the duodenum and jejunum after ingestion of food (36). The high degree of structural conservation of GIP among species supports the view that this peptide plays an important role in

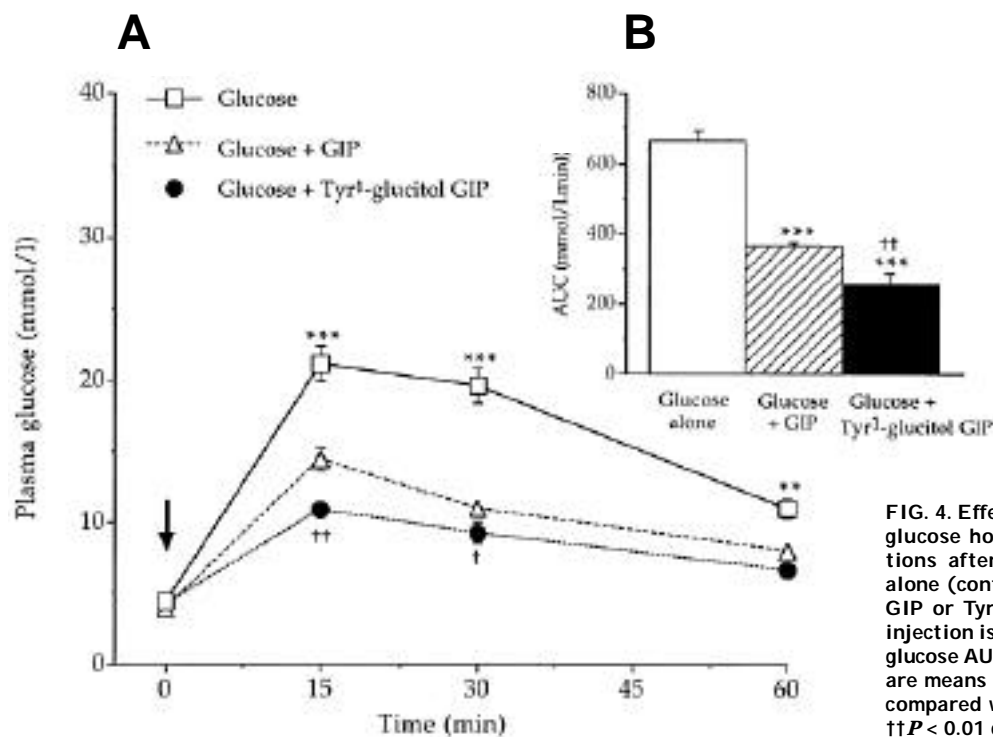


FIG. 4. Effects of GIP and Tyr¹-glucitol GIP on plasma glucose homeostasis. A: Plasma glucose concentrations after intraperitoneal glucose (18 mmol/kg), alone (control group), or in combination with either GIP or Tyr¹-glucitol GIP (10 nmol/kg). The time of injection is indicated by the arrow (0 min). B: Plasma glucose AUC values for 0–60 min postinjection. Values are means \pm SE for six rats. ** P < 0.01, *** P < 0.001 compared with GIP and Tyr¹-glucitol GIP; † P < 0.05, †† P < 0.01 compared with nonglycated GIP.

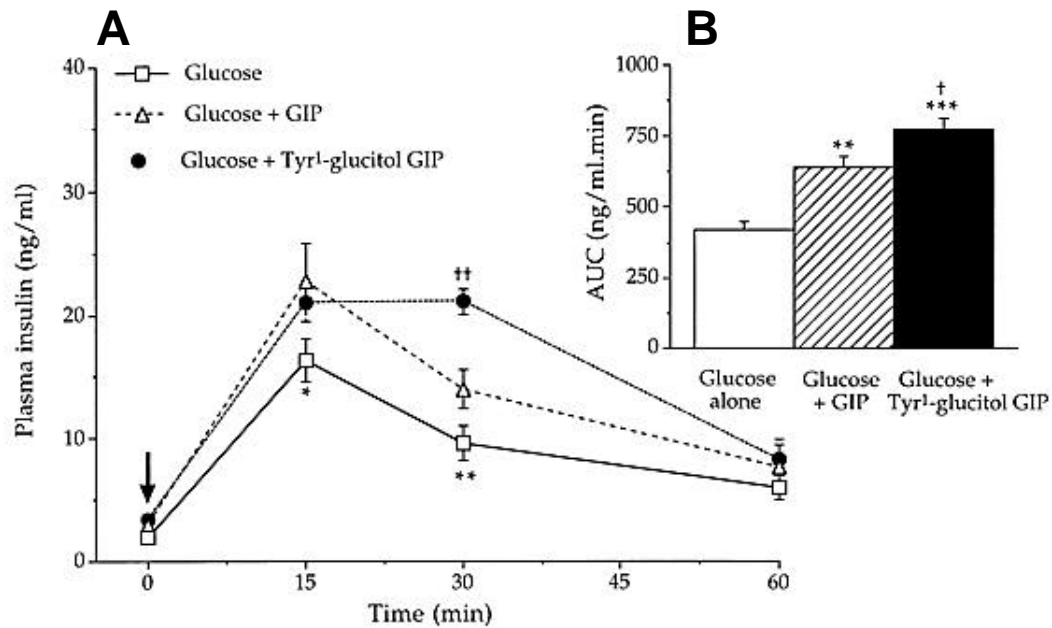


FIG. 5. Effects of GIP and Tyr¹-glucitol GIP on plasma insulin responses. **A:** Plasma insulin concentrations after intraperitoneal glucose (18 mmol/kg), alone (control group), or in combination with either with GIP or Tyr¹-glucitol GIP (10 nmol/kg). The time of injection is indicated by the arrow. **B:** Plasma insulin AUC values were calculated for each of the three groups up to 60 min postinjection. The time of injection is indicated by the arrow (0 min). Values are means \pm SE for six rats. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ compared with GIP and Tyr¹-glucitol GIP; † $P < 0.05$, †† $P < 0.01$ compared with nonglycated GIP.

metabolism (12). Secretion of GIP is stimulated directly by actively transported nutrients in the gut lumen without a notable input from autonomic nerves (12). The most important stimulants of GIP release are simple sugars (37) and unsaturated long-chain fatty acids (38), with amino acids exerting weaker effects (39). As with tGLP-1, the insulin-releasing effect of GIP is strictly glucose-dependent (30,40). This affords protection against hypoglycemia and thereby fulfills one of the most desirable features of any current or potential antidiabetic drug (41).

The present results demonstrate for the first time that Tyr¹-glucitol GIP displays profound resistance to serum and DPP IV degradation. Using ESI-MS, the present study showed that native GIP was rapidly cleaved in vitro to a major 4748.4-Da degradation product corresponding to GIP(3-42), which confirmed previous findings using matrix-assisted laser desorption ionization time-of-flight mass spectrometry (42). Serum degradation was completely inhibited by DPA (Ile-Pro-Ile), a specific competitive inhibitor of DPP IV, confirming this as the main enzyme for GIP inactivation in vivo (4,5). In contrast, Tyr¹-glucitol GIP remained almost completely intact after incubation with serum or DPP IV for up to 12 h. This indicates that glycation of GIP at the amino-terminal Tyr¹ residue masks the potential cleavage site from DPP IV and prevents removal of the Tyr¹-Ala² dipeptide from the NH₂-terminus, preventing the formation of GIP(3-42).

Consistent with in vitro protection against DPP IV, administration of Tyr¹-glucitol GIP significantly enhanced the antihyperglycemic activity and insulin-releasing action of the peptide when administered with glucose to rats. Native GIP enhanced insulin release and reduced the glycemic excursion as observed in many previous studies (12,40). However, amino-terminal glycation of GIP increased the insulin-releasing and antihyperglycemic actions of the peptide by 62 and

38%, respectively, as estimated from AUC measurements. Detailed kinetic analysis is difficult because of necessary limitation of sampling times, but the greater insulin concentrations after Tyr¹-glucitol GIP as opposed to GIP at 30 min postinjection is indicative of longer half-life. The glycemic rise was modest in both peptide-treated groups, and glucose concentrations after injection of Tyr¹-glucitol GIP were consistently lower than after GIP. Since the insulinotropic actions of GIP are glucose-dependent (30,40), it is likely that the relative insulin-releasing potency of Tyr¹-glucitol GIP is greatly underestimated in the present in vivo experiments.

In keeping with this interpretation, recent in vitro studies in our laboratory using glucose-responsive clonal β -cells showed that the insulin-releasing potency of Tyr¹-glucitol GIP was several orders of magnitude greater than that of GIP and that its effectiveness was more sensitive to change of glucose concentrations within the physiologic range (32). Together with the present in vivo observations, this suggests that NH₂-terminal glycation of GIP confers resistance to DPP IV degradation while enhancing receptor binding and insulin secretory effects on the β -cell. These attributes of Tyr¹-glucitol GIP are fully expressed in vivo, where DPP IV resistance impedes degradation of the peptide to GIP(3-42), thereby prolonging the half-life and enhancing effective concentrations of the intact biologically active peptide. It is thus possible that glycated GIP is enhancing insulin secretion in vivo by both enhanced potency at the receptor and improved DPP IV resistance. Thus numerous studies have shown that GIP(3-42) and other NH₂-terminally modified fragments, including GIP(4-42) and GIP(17-42), are either weakly effective or inactive in stimulating insulin release (4,43-45). Furthermore, evidence exists that NH₂-terminal deletions of GIP result in receptor antagonist properties in GIP receptor transfected Chinese hamster kidney cells (9), suggesting that inhibition of GIP catabolism would also reduce

the possible feedback antagonism at the receptor level by the truncated GIP(3-42).

In addition to its insulinotropic actions, a number of other potentially important extrapancreatic actions of GIP may contribute to the enhanced antihyperglycemic activity and other beneficial metabolic effects of Tyr¹-glucitol GIP. These include the stimulation of glucose uptake in adipocytes, increased synthesis of fatty acids, and activation of lipoprotein lipase in adipose tissue (46–48). GIP also promotes plasma triglyceride clearance in response to oral fat loading (49). In liver, GIP has been shown to enhance insulin-dependent inhibition of glycogenolysis (50). GIP also reduces glucagon-stimulated lipolysis in adipose tissue as well as hepatic glucose production (51). Finally, recent findings indicate that GIP has a potent effect on glucose uptake and metabolism in mouse isolated diaphragm muscle (52). The latter action may be shared with tGLP-1 (53,54), and both peptides have additional benefits of stimulating somatostatin secretion and slowing down gastric emptying and nutrient absorption (1,55).

In conclusion, this study has demonstrated for the first time that the glycation of GIP at the amino-terminal Tyr¹ residue limits GIP catabolism through impairment of the proteolytic actions of serum peptidases and thus prolongs its half-life in vivo. This effect is accompanied by enhanced antihyperglycemic activity and raised insulin concentrations in vivo, suggesting that such DPP IV-resistant analogs should be explored alongside tGLP-1 as potentially useful therapeutic agents for type 2 diabetes. Tyr¹-glucitol GIP appears to be particularly interesting in this regard, since such amino-terminal modification of GIP enhances (32) rather than impairs glucose-dependent insulinotropic potency, as was observed recently for tGLP-1 (28).

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