

Article

Inhibitory Potential of Synthetic Amino Acid Derivatives Against Digestive Enzymes—Promising Hypoglycemic and Anti-Obesity Agents

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Abstract: Over the last decades, the increased incidence of metabolic disorders such as type 2 diabetes and obesity has motivated researchers to investigate new enzyme inhibitors. In this study, the inhibitory effects of synthetic amino acid derivatives (PPC80, PPC82, PPC84, PPC89, and PPC101) on the activity of digestive enzymes was assessed by *in vitro* assays. The inhibitory effect was determined by the inhibition percentage and the 50% inhibitory concentration (IC₅₀), and the mechanism of action was investigated by Lineweaver–Burk plots. PPC80, PPC82, and PPC84 inhibited pancreatic lipase (IC₅₀ of 1.67–10.23 × 10⁻¹ mmol/L). The activity of pancreatic α-amylase was suppressed by PPC80, PPC82, PPC84, PPC89, and PPC101 (IC₅₀ of 1.62–5.19 × 10⁻¹ mmol/L). Finally, PPC84, PPC89, and PPC101 also showed a potent inhibitory effect on α-glucosidase (IC₅₀ of 0.51–3.53 × 10⁻¹ mmol/L). PPC80 and PPC82 followed a non-competitive inhibition mechanism against pancreatic lipase, while PPC84 acted through competitive inhibition. The inhibition of pancreatic α-amylase by the derivatives was non-competitive, as well as for PPC84, PPC89 and PPC101 against α-glucosidase. The results suggest that these synthetic amino acid derivatives have inhibitory potential against digestive enzymes and may be used as therapeutic agents to control metabolic disorders.

Keywords: Amino acid derivatives; Digestive enzymes; Pancreatic lipase; Pancreatic α-Amylase; α-Glucosidase; Metabolic disorders

1. Introduction

The digestion of foods in the gastrointestinal tract of humans and animals is determined by the activity of enzymes that break down macronutrients into smaller molecules to be absorbed in the gut and used by the body [1]. Some of the most important digestive enzymes include α-amylase and α-glucosidase, which degrade carbohydrates to obtain energy, and lipases, which catalyze the cleavage of triglycerides to produce free fatty acids and monoacylglycerol that either meet metabolic needs or are re-esterified and stored as triglycerides in adipose tissue [2]. Diets rich in carbohydrates can lead to hyperglycemia, which is associated with high insulin levels in the blood and increased uptake of nutrients, leading to the accumulation of adipose tissue and obesity [3]. On the other hand, high-fat diets are associated with abnormally high levels of circulating fatty acids and subsequent ectopic deposition in non-adipose tissues, as well as lipid accumulation in the liver, heart, endothelium, nervous system, pancreas, and skeletal muscle, thereby causing an imbalance in homeostatic mechanisms regulating metabolism [2,4]. This imbalance may lead to health complications such as

metabolic disorders (e.g., dyslipidemia, hypertension, type 2 diabetes), cancers, respiratory diseases, digestive problems, and osteoarthritis [5].

Regulation of nutrient absorption (e.g., carbohydrates) through the inhibition of digestive enzymes is an effective manner to control metabolism. For example, acarbose can inhibit the activity of α -amylase and α -glucosidase enzymes by reducing glucose absorption and decreasing insulin secretion in postprandial glycemia, establishing a glycemic control mechanism associated with reduced glycosylated hemoglobin. This class of enzymatic inhibitors is indicated in patients with adequate fasting blood glucose and elevated postprandial blood glucose levels. In patients with impaired glucose tolerance, enzymatic inhibitors have been associated with a marked reduction in cardiovascular events and no risk of adverse side effects, such as weight gain or hypoglycemia [6]. Therefore, the development of α -amylase and α -glucosidase inhibitors is increasingly recognized as a therapeutic strategy for patients with carbohydrate metabolic disorders, including postprandial hyperglycemia and type 2 diabetes mellitus [7,8].

Obesity is a complex disease that involves an abnormal or excessive accumulation of fat in the body and constitutes a public health problem worldwide [9]. The control and treatment of this pathology are mainly aimed at avoiding health complications, as well as increasing life expectancy [9]. Among the available drugs, lipase inhibitors (e.g., orlistat) act by reducing the absorption of monoacylglycerol and thus lead to weight loss [7,8,10,11]. However, new synthetic anti-obesity agents, which may bring better benefits to patients, have been investigated [12].

In this context, the preparation of novel amino acid derivatives obtained from organic synthesis processes is a promising area, which has been subjected to numerous biological studies. In addition to the functionalization of carboxylic and amine groups attached to the stereogenic center, the coupling of carbon side chains may also result in functional amino acid derivative drugs synthesized by conventional chemical reactions (i.e., acylation, alkylation, and amidation) [13]. These derivatives have attracted recent scientific interest due to their multiple biological properties [14,15]. For example, cationic antimicrobial peptides hold promise as new alternative antibiotics with the potential to inhibit multi-drug resistant bacteria [16].

In the present study, the inhibitory effects of synthetic amino acid derivatives on digestive enzymes were assessed using *in vitro* assays. This exploratory study may have predictive value for developing new therapeutic agents against metabolic disorders such as type 2 diabetes mellitus and obesity.

2. Materials and Methods

2.1. Synthesis of amino acid derivatives

The evaluated as amino acid derivatives, compounds PPC80 (342.52 g/mol), PPC82 (314.47 g/mol), PPC84 (287.40 g/mol), PPC89 (370.58 g/mol), and PPC101 (469.76 g/mol), were synthesized according to our previous report in the literature [17].

2.2 Chemicals

The drugs and reagents used in this study were as follows: porcine pancreatic lipase, 50 mmol/L tris-HCl buffer (pH 8.0), *p*-nitrophenol palmitate, Triton-X 100, orlistat, porcine pancreatic α -amylase, 50 mmol/L Tris-HCl (pH 7.0), α -Glucosidase, 100 mmol/L citrate-phosphate buffer (pH 7.0), acarbose, and *p*-nitrophenyl- α -D-glycopyranoside (Sigma-Aldrich® Co., St. Louis, MO, USA), while dimethylsulfoxide and starch (Loja Synth®, Diadema, SP, Brazil). Unless noted, all chemicals utilized in the synthetic protocol were acquired from Sigma-Aldrich® Co., St. Louis, MO, USA and used as received.

2.3. Inhibitory activity on digestive enzymes

2.3.1. Pancreatic lipase inhibition assay

The pancreatic lipase inhibition assay was performed according to Santos et al. [18] with some modifications. The porcine pancreatic lipase (10 g/L) was incubated in 50 mmol/L Tris-HCl buffer (pH 8.0) containing 10 mmol/L CaCl₂ and 25 mmol/L NaCl. The *p*-nitrophenol palmitate substrate (8 mmol/L) was dissolved in 0.5%, w/v Triton-X 100. PPC80, while PPC82, PPC84, PPC89, and PPC101 amino acid derivatives and orlistat were solubilized in dimethylsulfoxide (DMSO) prepared at increasing concentrations ranging from 0.5 to 13.92 × 10⁻¹ mmol/L. A total of 100 μL of enzyme solution, 50 μL of *p*-nitrophenol palmitate substrate, and 50 μL of amino acid derivative sample or orlistat were added to the microplate wells. Next, microplates were incubated at four different time intervals (10, 20, 30, and 40 min) in a water bath at 37 °C, and the reaction was stopped in an ice bath. All reactions were carried out in triplicate. The absorbance of the products was measured at 405 nm using a microplate reader (Thermoplate®, TP-Reader, Wuxi City, Jiangsu, China).

2.3.2. Pancreatic α-amylase inhibition assay

The pancreatic α-amylase inhibition assay was carried out according to Freitas et al. [19] with some modifications. The porcine pancreatic α-amylase (1 mg/mL) was incubated in 50 mmol/L Tris-HCl buffer (pH 7.0) containing 10 mmol/L CaCl₂ and 1% starch. PPC80, PPC82, PPC84, PPC89, and PPC101 amino acid derivatives and acarbose were solubilized in DMSO prepared at increasing concentrations ranging from 0.15 to 15.90 × 10⁻¹ mmol/L. A total of 50 μL enzyme solution, 50 μL substrate, and 50 μL amino acid derivative sample or acarbose were added to the microplate wells. Afterward, microplates were pre-incubated for 10 min in a water bath at 37°C. A total of 100 μL substrate was added at each well, and microplates were incubated at four different time intervals (10, 20, 30, and 40 min) in a water bath at 37°C. The reaction was stopped using an ice bath. All reactions were carried out in triplicate. The absorbance of the products was measured at 405 nm using a microplate reader (Thermoplate®, TP-Reader, Wuxi City, Jiangsu, China).

2.3.3. α-Glucosidase inhibition assay

The inhibitory effect against α-glucosidase was carried out according to Chelladurai and Chinnachamy [20] with some modifications. A total of 2 U/mL α-Glucosidase and 5 mmol/L *p*-nitrophenyl-α-D-glucopyranoside substrate were solubilized in 100 mmol/L citrate-phosphate buffer (pH 7.0). PPC80, PPC82, PPC84, PPC89, and PPC101 amino acid derivatives and acarbose were solubilized in DMSO prepared at increasing concentrations ranging from 0.24 to 17.40 × 10⁻¹ mmol/L. A total of 100 μL α-glucosidase solution, 50 μL amino acid derivative sample or acarbose, and 50 μL substrate were added to the microplate wells. Afterward, microplates were incubated at different intervals (10, 20, 30, and 40 min) in a water bath at 37°C. The reaction was stopped in an ice bath. All enzyme reactions were carried out in triplicate. The absorbance of the products was measured at 405 nm using a microplate reader (Thermoplate®, TP-Reader, Wuxi City, Jiangsu, China).

2.3.4. Determination of the inhibitory effect and IC₅₀

The percentage of inhibition (*I*%) was determined using 'absorbance versus time' graphs. By means of linear regression, using the method of least-squares, the equations of the straight lines and the angular coefficients were obtained to determine the inhibition (*I*%) of the enzymatic activities by the equation:

$$I\% = 100 \times \frac{(A - a) - (B - b)}{(A - a)}$$

where *A* is the angular coefficient of the straight-line equation (enzyme + substrate), *a* is the angular coefficient of the equation of the line (substrate), *B* is the angular coefficient of the straight-line equation (enzyme + substrate + sample), and *b* is the value of the angular coefficient of the straight-line equation (enzyme + sample).

The 50% inhibitory concentrations (IC_{50}) were determined through 'response *versus* concentration' plots using the linear least-squares regression model.

2.3.5. Determination of kinetic parameters

Kinetic parameters were determined using the same experimental conditions described above for each enzyme [21]. The reactions were prepared using increased substrate concentrations (16 to 0.06 mmol/L), both in the absence and presence of PPC80, PPC82, PPC84, PPC89, and PPC101 derivatives or positive control (orlistat or acarbose). The enzyme concentrations were maintained as described above. The absorbance of the products was measured at 405 nm using a microplate reader (Thermoplate®, TP-Reader, Wuxi City, Jiangsu, China) as a function of time (60 sec). The absorbance values were converted into product concentration ($\mu\text{mol/L}$) using standard curves of glucose (α -amylase) and *p*-nitrophenol (pancreatic lipase and α -glucosidase). The value of the initial velocity (v_0) of enzymatic reactions was estimated to create the ' v_0 *versus* substrate concentration' graph. Kinetic constants (K_m or K_i and V_{max}) were calculated, and the inhibition model was verified using Lineweaver–Burk plots [21].

2.4. Statistical analysis

The data were subject to the analysis of variance (ANOVA) and Tukey's test ($p < 0.05$) to determine the differences between mean groups using the GraphPad Prism 5 program. Data were presented as mean \pm S.E.M.

3. Results

3.1. Synthesis of protected amino acid derivatives

The synthesis started by reacting Boc-protected L-isoleucine amino acid with EDC.HCl as carboxylic acid coupling. After 30 min, the vessel was charged with the corresponding nucleophile in the presence of racemic camphorsulphonic acid (+/-)-CSA as organocatalyst [17]. The corresponding synthetic amino acid derivatives PPC80, PPC82, PPC84, PPC89, and PPC101 were attaining in yields ranging from 67 to 80% (Figure 1). It is worth to mention, no epimerization process was observed. Characterization data are in agreement with those previously described in the literature [17]. The products were then used to carry out inhibitory activity assays against digestive enzymes.

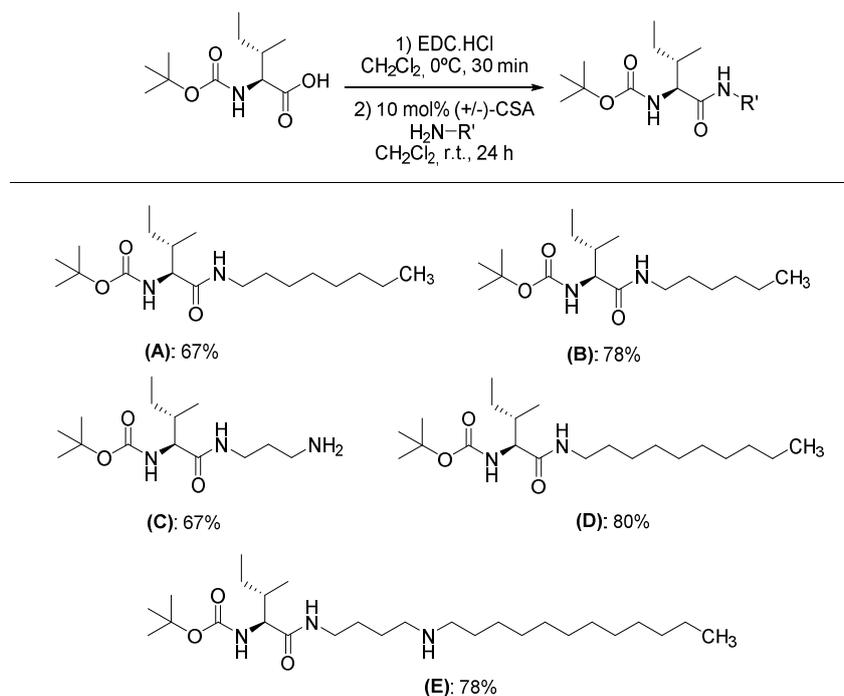


Figure 1. Synthesis and molecular structure of amino acid derivatives. (A) PPC 80; (B) PPC 82; (C) PPC 84; (D) PPC 89; (E) PPC 101.

3.2. Inhibitory effect of amino acid derivatives on pancreatic lipase

The results showed that the inhibitory effect of PPC82, PPC80, and PPC84 amino acid derivatives on pancreatic lipase activity was concentration-dependent (Figure 2). PPC80 (5.84×10^{-1} mmol/L) was more active ($p < 0.05$) than orlistat (8.07×10^{-1} mmol/L, 60% inhibition) at a lower concentration, showing an inhibitory effect of about 65% on pancreatic lipase activity (Figure 2A). Also, PPC82 (4.77×10^{-1} mmol/L) was more effective in inhibiting pancreatic lipase at a lower concentration than orlistat (8.07×10^{-1} mmol/L), with a response of about 86% (Figure 2B). Moreover, PPC84 (13.92×10^{-1} mmol/L) exerted a similar inhibitory effect on pancreatic lipase at a higher concentration than orlistat (8.07×10^{-1} mmol/L), showing an inhibitory effect of about 62% (Figure 2C). In this assay, PPC89 and PPC101 did not show inhibitory action on the reference enzyme.

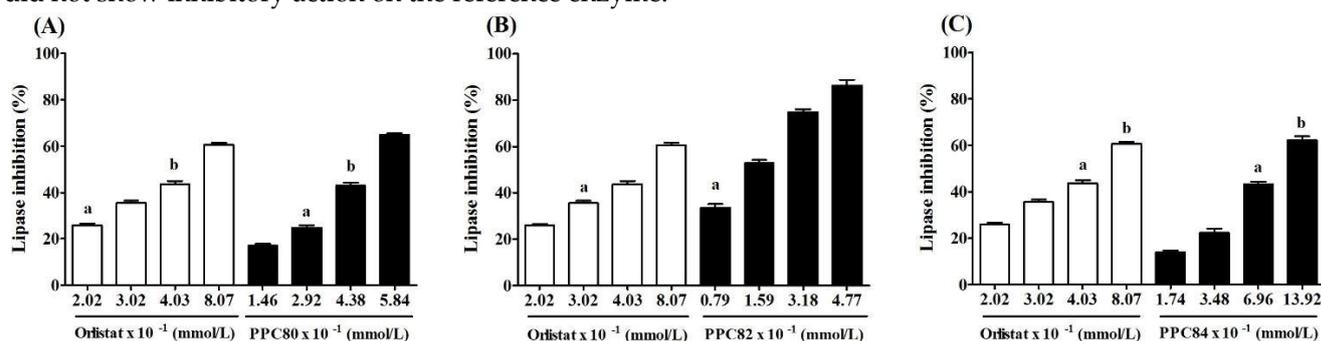


Figure 2. Inhibitory effects of amino acid derivatives on pancreatic lipase activity. Each bar represents the mean \pm S.E.M. ($n = 3$). Repeated letters in the same figure, group means did not show statistically significant differences after ANOVA and Tukey's test ($p < 0.05$).

As shown in Table 1, the IC_{50} values were also determined. PPC80 and PPC82 showed better IC_{50} values than orlistat at the lower concentrations of 4.75 ± 0.08 , 1.67 ± 0.06 , and $5.88 \pm 0.15 \times 10^{-1}$ mmol/L, respectively ($p < 0.05$). On the contrary, PPC84 ($10.23 \pm 0.20 \times 10^{-1}$ mmol/L) had a higher IC_{50} value at a lower concentration than orlistat ($p < 0.05$), thereby showing a lower inhibitory effect on pancreatic lipase activity. As noted in Figure 2C, PPC84 concentration was almost 2-fold higher than the

reference compound, which may be associated with its high concentration (13.92×10^{-1} mmol/L) observed in the inhibitory effect assay (Figure 2C) and shows a low inhibitory affinity with the target enzyme.

Table 1. IC₅₀ values of amino acid derivatives against pancreatic lipase, pancreatic α -amylase and α -glucosidase.

Group	IC ₅₀ ($\times 10^{-1}$ mmol/L)		
	pancreatic lipase	pancreatic α -amylase	α -glucosidase
Orlistat	5.88 \pm 0.15	-	-
Acarbose	-	3.26 \pm 0.03 ^a	6.39 \pm 0.05
PPC80	4.75 \pm 0.08	2.76 \pm 0.05 ^a	-
PPC82	1.67 \pm 0.06	5.19 \pm 0.20 ^b	-
PPC84	10.23 \pm 0.20	4.93 \pm 0.11 ^b	3.21 \pm 0.02
PPC89	-	1.71 \pm 0.14 ^c	3.53 \pm 0.06
PPC101	-	1.62 \pm 0.02 ^c	0.51 \pm 0.02

Each value represents the mean \pm S.E.M (n = 3). Repeated letters in the same column, group means did not show statistically significant differences after ANOVA and Tukey's test (p < 0.05).

3.3. Kinetic parameters on pancreatic lipase activity

The inhibitory mechanism of PPC80, PPC82, and PPC84 compounds on the pancreatic lipase activity was assessed. The Lineweaver–Burk profiles showed that PPC80 and PPC82 follow a non-competitive inhibition mechanism (Figure 3A-C). As observed in the Lineweaver–Burk plots, the straight lines did intersect the x-axis but did not the y-axis, thereby indicating a decrease in the maximum velocity (V_{max}) of the reaction [21]. On the other hand, PPC84 followed a competitive inhibition mechanism since an intersection on the y-axis was observed (Figure 3). These profiles were used to determine the values of kinetic parameters (Table 2).

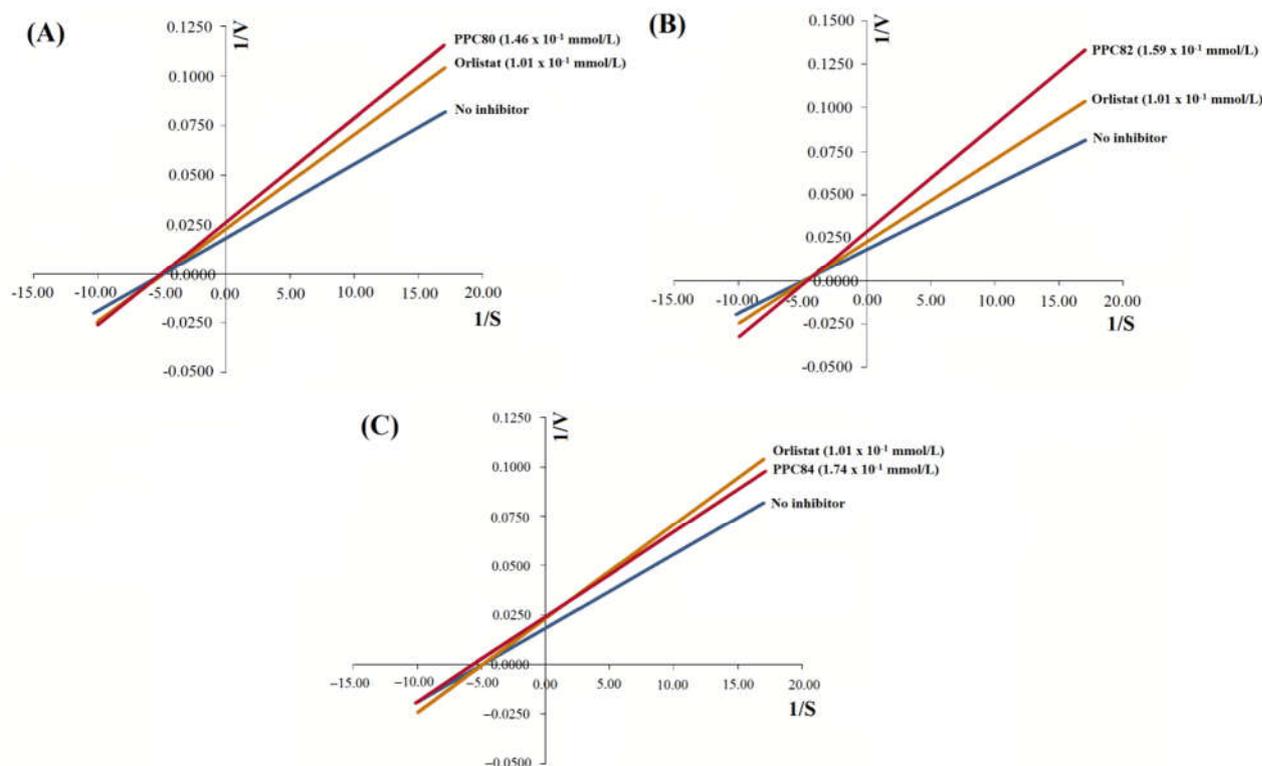


Figure 3. Lineweaver-Burk kinetic profile of amino acid derivatives against pancreatic lipase.

Table 2. Kinetic parameters of PPC80, PPC82 and PPC84 against pancreatic lipase.

Group	Concentration (mmol/L)	K_m/K_i (mmol/L)	V_{max} ($\mu\text{mol/L}\cdot\text{min}$)
No inhibitor	-	0.204 ± 0.003^a	55.78 ± 2.05
Orlistate	1.01×10^{-1}	0.209 ± 0.001^{ab}	43.75 ± 0.76^a
PPC80	1.46×10^{-1}	0.211 ± 0.004^{abc}	41.36 ± 0.05^{ab}
PPC82	1.59×10^{-1}	0.221 ± 0.001^c	35.05 ± 0.13
PPC84	1.74×10^{-1}	0.181 ± 0.003	41.36 ± 0.52^{ab}

Each value represents the mean \pm S.E.M (n = 3). Repeated letters in the same column, group means did not show statistically significant differences after ANOVA and Tukey's test ($p < 0.05$).

The kinetic parameters of the enzyme-substrate reaction in the absence of enzyme inhibitors were $K_m = 0.204 \pm 0.003$ mmol/L and $V_{max} = 55.78 \pm 2.05$ $\mu\text{mol}/\text{min}$ per L (Table 2). The addition of 1.01 mmol/L orlistat to the reaction caused a significant decrease in V_{max} (43.75 ± 0.76 $\mu\text{mol}/\text{min}$ per L; $p < 0.05$) and an increase in K_i (0.209 ± 0.001 mmol/L; $p < 0.05$), indicating that the reaction rate became slower in the presence of this compound. Moreover, the addition of 1.46 mmol/L PPC80 to the reaction reduced the reaction rate, causing an increase in K_i (0.211 ± 0.004 mmol/L) and a decrease in V_{max} (41.36 ± 0.05 $\mu\text{mol}/\text{min}$ per L), while 1.59 mmol/L PPC82 was more active ($K_i = 0.221 \pm 0.001$ mmol/L; $V_{max} = 35.05 \pm 0.13$ $\mu\text{mol}/\text{min}$ per L). Taken together, these results and those shown in Supplementary materials 1A and 1B suggest that PPC80 and PPC82 exert a non-competitive inhibitory effect on pancreatic lipase. Moreover, 1.74 mmol/L PPC84 reduced V_{max} (41.36 ± 0.52 $\mu\text{mol}/\text{min}$ per L) and K_i (0.181 ± 0.003 mmol/L), showing that this compound exerted competitive inhibition as observed in the Lineweaver-Burk plot (Figure 3C).

3.4. Inhibitory effect of amino acid derivatives on pancreatic α -amylase

The results showed that the inhibitory effect of PPC80, PPC82, PPC84, PPC89, and PPC101 amino acid derivatives on pancreatic α -amylase activity was concentration-dependent (Figure 4). PPC80 (7.30×10^{-1} mmol/L), PPC82 (15.90×10^{-1} mmol/L), PPC84 (8.70×10^{-1} mmol/L), PPC89 (6.75×10^{-1} mmol/L), and PPC101 (5.32×10^{-1} mmol/L) inhibited pancreatic α -amylase by nearly 93%, 94%, 74%, 90%, and 86% ($p < 0.05$), respectively, while acarbose (6.20×10^{-1} mmol/L) reduced the specific enzymatic activity by about 86% (Figures 4A–E). PPC101 was more active than acarbose at a lower concentration, while PPC80 and PPC82 showed better inhibitory effects than acarbose at higher concentrations. Moreover, PPC89 (4.05×10^{-1} mmol/L) produced the same inhibitory effect as acarbose (6.20×10^{-1} mmol/L).

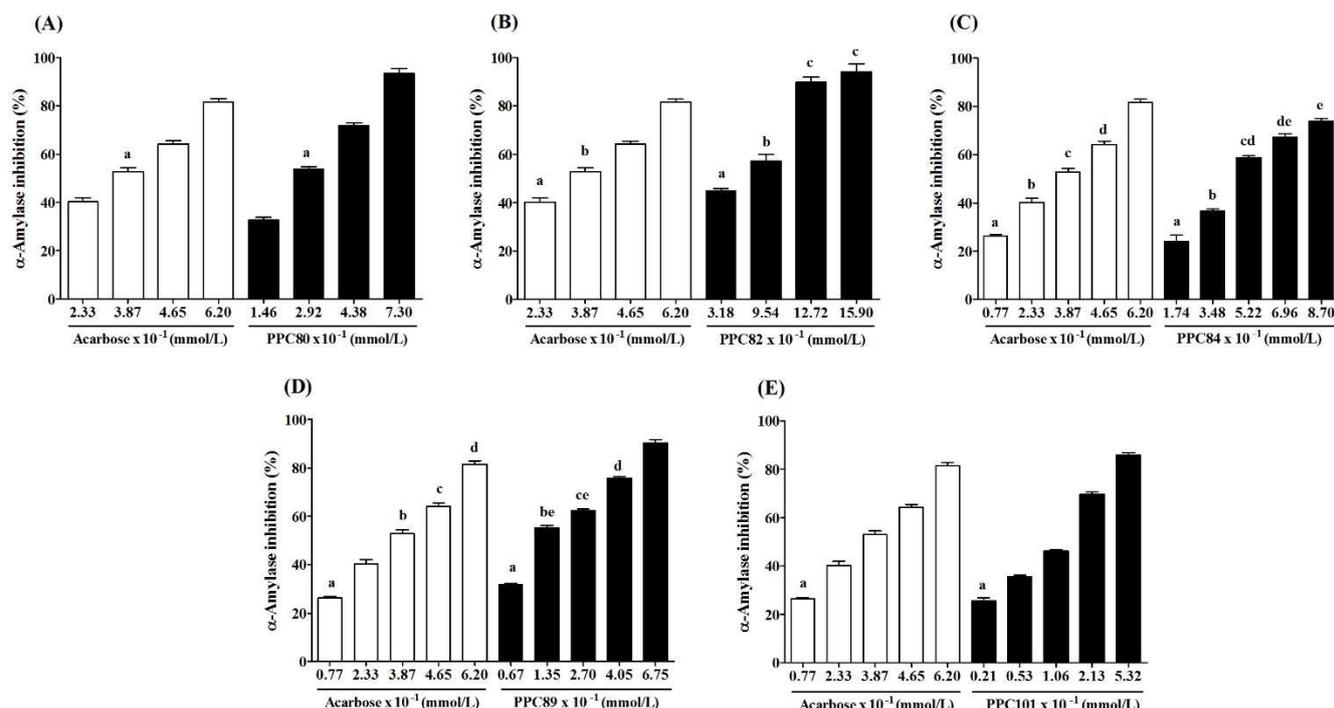


Figure 4. Inhibitory effect of amino acid derivatives on pancreatic α -amylase. Each bar represents mean \pm S.E.M. ($n = 3$). Repeated letters in the same figure, group means did not show statistically significant differences after ANOVA and Tukey's test ($p < 0.05$).

The IC_{50} values showed the inhibitory potential of PPC80, PPC82, PPC84, PPC89 and PPC1010 derivatives (Table 1). PPC89 ($1.71 \pm 0.14 \times 10^{-1}$ mmol/L) and PPC101 ($1.62 \pm 0.02 \times 10^{-1}$ mmol/L) had lower IC_{50} values than acarbose ($3.26 \pm 0.03 \times 10^{-1}$ mmol/L), thereby showing a more potent inhibitory activity against pancreatic amylase, while PPC82 and PPC84 were less effective in inhibiting the activity of the enzyme ($p < 0.05$). Moreover, PPC80 showed a similar suppressive potential as the reference compound (acarbose). The inhibitory effects were corroborated in Figure 4, where the inhibition of the pancreatic α -amylase enzyme occurred at higher concentrations of PPC82 and PPC84.

3.5. Kinetic parameters against pancreatic α -amylase

The kinetic parameters of pancreatic α -amylase activity for PPC89, PPC101, PPC80, PPC84, and PPC82 amino acid derivatives were determined. The Lineweaver–Burk plots showed that the inhibition mechanism of PPC80, PPC82, PPC84, PPC89, and PPC101 is non-competitive (Figure 5A–B), as a single point of intersection on the x-axis was verified [21]. These profiles were then used to determine the values of kinetic parameters (Table 3).

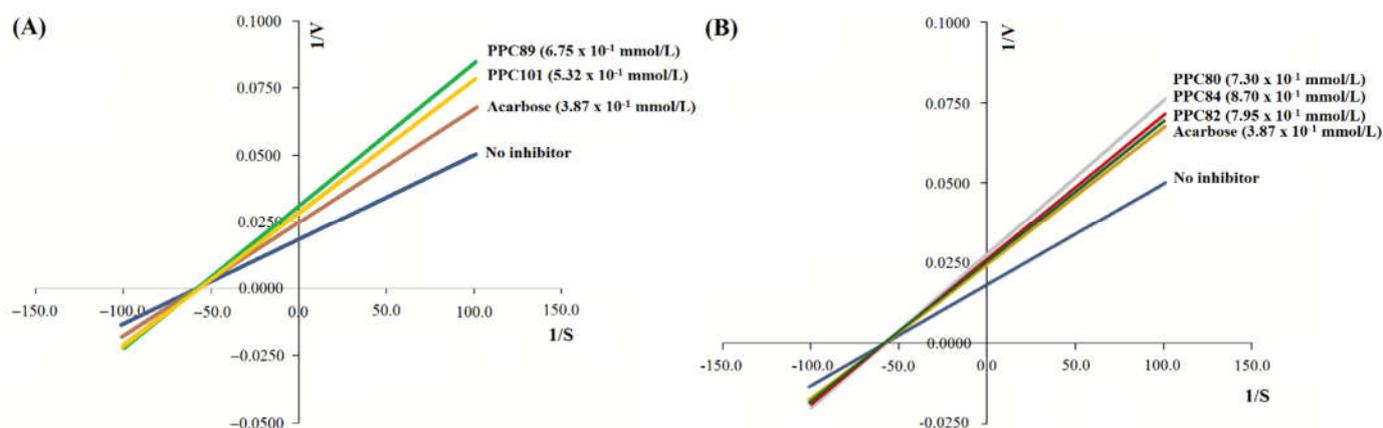


Figure 5. Lineweaver-Burk kinetic profile of amino acid derivatives against pancreatic amylase.

Table 3. Kinetic parameters of amino acid derivatives against pancreatic α -amylase.

Group	Concentration (mmol/L)	K_m/K_i (mmol/L)	V_{max} ($\mu\text{mol/L}\cdot\text{min}$)
No inhibitor	-	0.153 ± 0.006	102.06 ± 0.77
Acarbose	3.87×10^{-1}	0.232 ± 0.002^a	71.78 ± 0.76^a
PPC80	7.30×10^{-1}	0.258 ± 0.004^a	63.72 ± 0.69
PPC82	7.95×10^{-1}	0.238 ± 0.008^{ab}	70.69 ± 0.54^{ab}
PPC84	8.70×10^{-1}	0.219 ± 0.007^{ab}	68.36 ± 0.92^{ab}
PPC89	6.75×10^{-1}	0.312 ± 0.005^c	57.35 ± 1.09^c
PPC101	5.32×10^{-1}	0.294 ± 0.004^c	59.05 ± 0.61^c

Each value represents the mean \pm S.E.M ($n = 3$). Repeated letters in the same column, group means did not show statistically significant differences after ANOVA and Tukey's test ($p < 0.05$).

In the absence of enzyme inhibitors, the reaction had a lower K_m (0.153 ± 0.006 mmol/L) and a higher V_{max} (102.06 ± 0.77 $\mu\text{mol}/\text{min}$ per L) than in the presence of acarbose or amino acid derivatives ($p < 0.05$), showing that these compounds can inhibit pancreatic amylase and reduce the reaction rate (Table 3). PPC80, PPC82, PPC84, PPC89, and PPC101 derivatives decreased V_{max} in a range from 57.35 ± 1.09 $\mu\text{mol}/\text{min}$ per L to 71.78 ± 0.78 $\mu\text{mol}/\text{min}$ per L, and inhibition constant (K_i) values ranged from 0.219 ± 0.007 mmol/L to 0.312 ± 0.005 mmol/L. The kinetic data and Lineweaver–Burk plots (Figure 5A-B) suggested that amino acid derivatives inhibit enzymatic function through a non-competitive inhibitory mechanism. In line with these results, a previous study also reported non-competitive inhibition for synthetic peptides that showed V_{max} values ranging from 0.313 and 0.558 [22].

3.6. Inhibitory effect of amino acid derivatives on α -glucosidase

As shown in Figure 6, the inhibitory effect of PPC84, PPC89, and PPC101 amino acid derivatives on α -glucosidase activity was concentration-dependent. PPC84 (4.35×10^{-1} mmol/L), PPC89 (6.74×10^{-1} mmol/L), and PPC101 (0.67×10^{-1} mmol/L) inhibited α -glucosidase by nearly 66, 78 and 64% ($p < 0.05$), respectively, inhibiting enzyme activity at lower concentrations than acarbose (positive control). These derivatives were also more effective at higher concentrations (Figure 6A-C). Moreover, PPC80 and PPC82 did not show inhibitory action against the tested enzyme.

The IC_{50} values showed the inhibitory potential of PPC84, PPC89, and PPC101 amino acid derivatives on α -glucosidase activity (Table 1). PPC89, PPC84, and PPC101 derivatives had lower IC_{50} and therefore were more effective for inhibiting α -glucosidase than acarbose ($IC_{50} = 6.39 \pm 0.05 \times 10^{-1}$ mmol/L). It is worth noting that PPC101 was 12-fold more potent than acarbose, exhibiting an outstanding potential to inhibit the target enzyme.

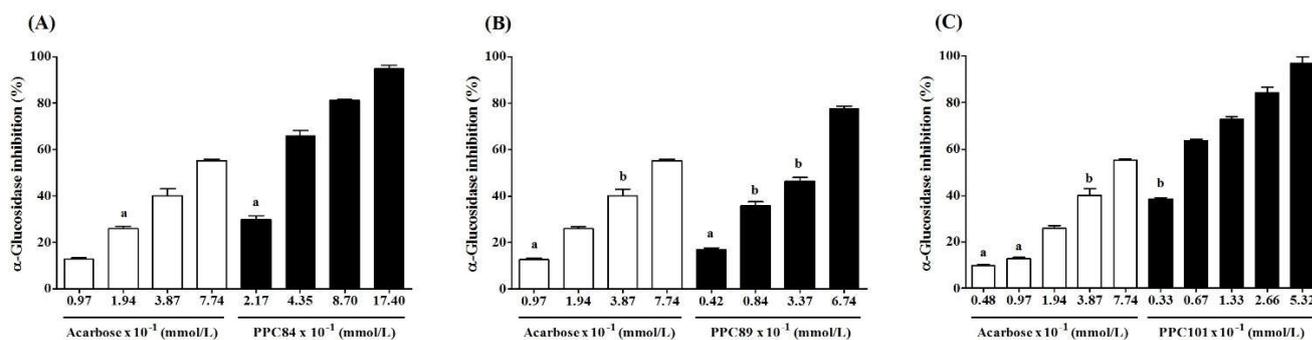


Figure 6. Inhibitory effect of amino acid derivatives on α -glucosidase. Each bar represents mean \pm S.E.M. (n = 3). Repeated letters in the same figure, group means did not show statistically significant differences after ANOVA and Tukey's test ($p < 0.05$).

3.7. Kinetic parameters of α -glucosidase

The inhibition assay and the Lineweaver–Burk plot (Figure 7) revealed that the inhibition mechanism of PPC101, PPC89, and PPC84 against α -glucosidase is non-competitive. This mechanism is related to the crossing of inhibition lines at an intersection point on the x-axis [21]. These data were then used to determine the values of kinetic parameters (Table 4).

In the absence of enzyme inhibitors, the reaction was faster than acarbose and had a K_m of 0.184 ± 0.006 mmol/L and a V_{max} of 78.26 ± 0.09 μ mol/min per L (Table 4). The presence of acarbose reduced the reaction rate, showing K_i and V_{max} values of 0.209 ± 0.006 mmol/L and 56.64 ± 0.15 μ mol/min per L, respectively. PPC84, PPC89, and PPC101 showed significantly lower V_{max} and higher K_i values ($p < 0.05$) than the reaction without the inhibitor. Therefore, PPC84, PPC89, and PPC101 derivatives reduced the reaction rate and confirmed the results observed in the Lineweaver–Burk plots (Figure 7).

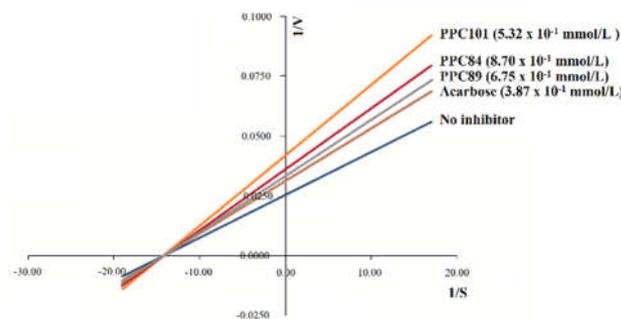


Figure 7. Lineweaver-Burk kinetic profile of amino acid derivatives against α -glucosidase.

Table 4. Kinetic parameters of amino acid derivatives against α -glucosidase.

Group	Concentration (mmol/L)	K_m/K_i (mmol/L)	V_{max} (μ mol/L.min)
No inhibitor	-	0.184 ± 0.006	78.26 ± 0.09
Acarbose	3.87×10^{-1}	0.209 ± 0.007^a	56.64 ± 0.15
PPC84	8.70×10^{-1}	0.237 ± 0.001	48.05 ± 0.06
PPC89	6.75×10^{-1}	0.214 ± 0.001^a	53.72 ± 0.02
PPC101	5.32×10^{-1}	0.269 ± 0.001	42.37 ± 0.03

Each value represents the mean \pm S.E.M (n = 3). Repeated letters in the same column, group means did not show statistically significant differences after ANOVA and Tukey's test ($p < 0.05$).

4. Discussion

The structure of amino acid derivatives suggested that the hydrocarbon chain is involved in their inhibitory effects since compounds with a side chain with more than eight carbon atoms did not inhibit pancreatic lipase. The inclusion of an amino group at the carbon side chain of PPC84 (Figure 1C) may have led to additional hydrogen bonding interactions (non-covalent interactions) at the catalytic site, resulting in an impaired ability to competitively inhibit the enzymatic activity. Besides, the possibility of the amine to act as a nucleophile (covalent bonding) cannot be ruled out [23].

These findings are consistent with previous studies assessing the effects of amino acid and peptide derivatives on pancreatic lipase activity. Ngoh and Gan [24] identified different peptides from common bean (*Phaseolus vulgaris*) that inhibited pancreatic lipase in the range of 23–87%. Polylysine is a synthetic peptide that also acts as a lipase inhibitor, showing a remarkable inhibition (80%) on the activity of porcine pancreatic lipase at a concentration of 100 mg/mL [25]. Furthermore, synthetic peptides [26] and hydrolyzed peptides [23] inhibited pancreatic lipases with IC₅₀ values below 50 μM.

The results of the present study showed that PPC80 and PPC82 have inhibitory potential on the activity of pancreatic lipase, which may promote a reduction of intestinal fat absorption and potentially affect body weight [27]. Therefore, PPC80 and PPC82 derivatives constitute promising therapeutic agents for the treatment of obesity and related lipid disorders.

Unlike competitive inhibitors, non-competitive inhibitors are unaffected by the substrate concentration, and therefore high concentrations of the inhibitor are not required to compete with the substrate [28]. Non-competitive inhibitors do not bind to the active site of the enzyme, but at another region called the allosteric site. Thus, non-competitive inhibitors can interact with the free enzyme to form the enzyme-inhibitor complex or with potential binding sites of the ES complex to form the enzyme-substrate-inhibitor complex, both of which are catalytically inactive. The formation of these complexes induces structural changes in the enzyme, modifying the conformation of the active site to prevent its interaction with the substrate [29].

The function of amino acid derivatives may be associated with the size of the hydrocarbon chain since compounds such as PPC89 and PPC101, which exhibit a high number of carbon atoms after nitrogen in their aliphatic chains, were more effective in inhibiting the activity of α-amylase at low IC₅₀ values. PPC80 (eight carbons) showed a similar inhibitory effect as acarbose, thus being the third most effective compound. Moreover, PPC82 (six carbons) and PPC84 (three carbons and one amino group) showed the lowest inhibitory activities.

The catalytic mechanism of the α-amylase family is stable and specific because of the α-retaining double displacement reaction. This two-step mechanism is a distinctive feature of the α-amylase family and may contribute to its broad specificity due to the attachment of different domains to the catalytic site or to extra sugar-binding subsites around the catalytic site [30]. However, the carboxylic groups of aspartate and glutamate residues can act as acid/base catalysts and nucleophilic reagents during the formation of covalent intermediates in the catalytic cycle. The presence of Cl⁻ ions may lead to the activation and facilitates the protonation of a carboxyl group [31].

An increasing number of studies have shown that synthetic compounds derived from amino acids and peptides exhibit an inhibitory action on α-amylase [22,32,33]. Admassu et al. [22] identified two α-amylase inhibitor peptides (GGSK and ELS) released by hydrolysis of proteolytic enzymes obtained from the red seaweed (*Porphyra* species). The IC₅₀ values for GGSK and ELS were 2.58 mM ± 0.08 and 2.62 mM ± 0.05, respectively. Another study reported peptides extracted from basil (*Ocimum basilicum*) seeds that showed 36% inhibition on α-amylase [33]. Similarly, González-Montoya et al. [32] also identified peptides from soy (*Glycine max*) protein capable of inhibiting pancreatic α-amylase activity at IC₅₀ values ranging from 0.16 to 8.30 mg/mL.

Pancreatic α-amylase and α-glucosidase are critical enzymes involved in the digestion of dietary starch, catalyzing the release of oligosaccharides that are further degraded into glucose. Therapeutic approaches for the treatment of type 2 diabetes include the inhibition of these enzymes to decrease the absorption of glucose in the digestive tract and reduce postprandial hyperglycemia [34,35]. Acarbose, miglitol, and voglibose are major inhibitors that reduce the rate of glucose absorption,

attenuating the postprandial increase in plasma glucose level and thus helping in the treatment of obesity [36,37]. Our results indicate that amino acid derivatives are potent inhibitors of pancreatic α -amylase and can be promising agents for the treatment of diabetes and metabolic disorders.

Several studies have reported the promising potential of amino acid and peptide derivatives as α -glucosidase inhibitors [37-39]. For example, KLPGF and NVLQPS peptides obtained from albumin showed inhibitory activity on α -glucosidase at IC_{50} values of $59.5 \pm 5.7 \mu\text{M}$ and $100.0 \pm 5.7 \mu\text{M}$, respectively [39]. In this study, the inhibitory activity of the KLPGF peptide motif was similar to acarbose ($IC_{50} = 60.8 \mu\text{M}$). Furthermore, three peptides isolated from quinoa (*Chenopodium quinoa*) showed similar inhibitory activity against α -glucosidase [37]. Singh and Kaur [38] reported serine-threonine-tyrosine-valine-containing peptides isolated from the endophytic fungi *Acacia nilotica* that exhibited potent inhibitory effects against α -glucosidase at low IC_{50} values ($3.75 \mu\text{g/mL}$).

The human α -glucosidase is an enzyme found in the epithelium of the small intestine that catalyzes starch breakdown and the consequent release of glucose. Therefore, inhibition of this enzyme constitutes a promising strategy to reduce serum glucose levels in metabolic diseases, including type 2 diabetes [20]. Our results showed that PPC89, PPC84, and PPC101 amino acid derivatives inhibit α -glucosidase, exhibiting potential as agents for lowering blood glucose levels in carbohydrate-related metabolic diseases.

Our results are in line with the reported inhibition mechanism for F5-SP, a peptide derivative with a K_i value of $86.63 \pm 0.014 \text{ ng/mL}$ that could bind to multiple sites of the α -glucosidase enzyme to modify its conformation [40].

5. Conclusions

In summary the results showed that PPC80, PPC82, PPC84, PPC89, and PPC101 amino acid derivatives are potential inhibitors of lipase, α -amylase, and α -glucosidase enzymes. For instance, PPC80, PPC82, and PPC84 inhibited pancreatic lipase, with IC_{50} values as low as 1.67 mmol/L . The activity of pancreatic α -amylase was suppressed by PPC80, PPC82, PPC84, PPC89, and PPC101, with IC_{50} values in a range of 1.62 – 5.19 mmol/L . In addition, PPC84, PPC89, and PPC101 also presented an inhibitory effect on α -glucosidase, with IC_{50} values as low as 0.51 mmol/L . PPC80 and PPC82 followed a non-competitive inhibition mechanism against pancreatic lipase, while PPC84 acted through competitive inhibition. Moreover, PPC80 and PPC82 followed a non-competitive inhibition mechanism against pancreatic lipase, while PPC94 acted through competitive inhibition. PPC80, PPC82, PPC84, PPC89, and PPC101 displayed non-competitive inhibition against α -amylase, while PPC84, PPC89, and PPC101 against α -glucosidase. The present study supports that amino acid derivatives are promising therapeutic agents for metabolic disorders, including type II diabetes and obesity.

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