

TRYPsin FROM JUMBO SQUID (*Dosidicus gigas*) HEPATOPANCREAS: PURIFICATION AND CHARACTERIZATION

Ana Gloria Villalba-Villalba, Enrique Márquez-Ríos, Marina Josafat Esquerra-Brauer and Francisco Javier Castillo-Yáñez

SUMMARY

Trypsin was purified from the hepatopancreas of *Dosidicus gigas* by fractionation with ammonium sulfate (30-70% saturation), gel filtration, affinity and ion exchange chromatography. The molecular weight of the trypsin obtained was ~23.5kDa according to sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and, showed a single band in the zymogram. The enzyme exhibited maximal activity at pH 8.5 and 40°C, using BAPNA as substrate. It was effectively inhibited by phenyl

methyl sulfonyl fluoride (PMSF) (98%), and *N*-*p*-Tosyl-L-lysine chloromethyl ketone (TLCK) (99%). Enzyme activity was affected by the following ions in decreasing order: Hg²⁺, Fe²⁺, Cu²⁺, Li⁺, Mg²⁺, K⁺, Mn²⁺, Ca²⁺. Trypsin activity decreased continuously as NaCl concentration increased from 0 to 30%. Km and kcat values were 0.085 ±1.45mM and 1.76 ±0.12s⁻¹, respectively. The results suggest that the purified enzyme a potential agent to be used in biotechnological processes.

Introduction

The marine environment contains the largest pool of diversified genetic material and, hence, represents an enormous potential source of enzymes. Among other enzymes, the possible applications of serine proteases in industry have been studied (Klomklao *et al.*, 2005).

The demand for enzymes with specific properties is large, and various enzymes sources are currently being investigated. It has been reported that proteases of marine sources exhibit high catalytic activity (Shahidi *et al.*, 2001). Jumbo squid is an important fishery in Mexico, usually sold as frozen or cooked-frozen gutted mantle (Luna-Raya *et al.*, 2006), after the hepatopancreas is removed, as it represents a source of contamination. This organ could be, however, a good source of enzymes. Jumbo

squid hepatopancreas constitutes 2-8% of the body weight and contains 18-20% crude protein (Esquerra-Brauer *et al.*, 2002). Extracts of hepatopancreas from the Short finned squid, *Illex illecebrosus*, have been experimentally used in a number of food processing applications, including the preparation of capelin fish sauce; salted, fermented squid mantle; squid tenderization and cheddar cheese ripening (Lee *et al.*, 1982). Fish sauce, with squid hepatopancreas as a fermentation aid, has been commercially produced in Canada for more than 10 years (Raksakulthai and Haard, 1999).

Included in the five classes of proteinases catalyzing the hydrolysis of peptide bonds, serine endoproteinases are divided into two superfamilies that evolved independently a similar catalytic mechanism. The trypsin superfamily includes trypsins and chymotryp-

sins, which are ubiquitous in animals (Hu and Leger, 2014). One of the most important fish and aquatic invertebrate viscera digestive enzyme is trypsin (EC 3.4.21.4) because they take part in a number of physiological processes, playing a major and well understood key role in hydrolyzing food protein (Diaz-Mendoza *et al.*, 2005; Delcroix *et al.*, 2006). This enzyme cleaves the peptide bond on the carboxyl side of arginine and lysine, has endopeptidase activity and a molecular weight ranging from 22 to 28kDa (Whitaker, 1994). The first studies of proteinases, mainly trypsin and chymotrypsin in vertebrate species, were carried out during the first part of the 20th century, describing kinetic properties, isolation and some molecular characteristics (Muhlía-Almazán *et al.*, 2008). Trypsin has been isolated and characterized thoroughly based on their physical-

chemical and enzymatic properties from several aquatic species, such as *Siniperca chuatsi* (Lu *et al.*, 2008), *Balistes capricus* (Jellouli *et al.*, 2009), *Sepia officinalis* (Balti *et al.*, 2009), *Lutjanus vitta* (Khantaphant and Benjakul, 2010), *Diapterus rhombeus* (Silva *et al.*, 2011), *Salaria basilisca* (Ktari *et al.*, 2012) and *Pterygoplichthys disjunctivus* (Villalba-Villalba *et al.*, 2013). However, most studies have been done in fishes, with very scarce information on trypsin from cephalopods. The aim of this study was to purify and characterize trypsin from the jumbo squid (*Dosidicus gigas*) hepatopancreas, thus generating basic information about this by-product.

Materials and Methods

Jumbo squid samples

Jumbo squid specimens were captured in the Gulf of

KEYWORDS / *Dosidicus gigas* / Enzyme Purification / Hepatopancreas / Squid / Trypsin /

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TRIPSINA DEL HEPATOPANCREAS DE CALAMAR GIGANTE (*Dosidicus Gigas*): PURIFICACIÓN Y CARACTERIZACIÓN

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RESUMEN

Se logró purificar tripsina del hepatopancreas de *Dosidicus gigas* por fraccionación con sulfato de amonio (30-70% de saturación), cromatografía de filtración en gel, afinidad e intercambio iónico. El peso molecular de la tripsina fue de 23,5 kDa, según la electroforesis SDS-PAGE y mostró una sola banda en el zimograma. La enzima mostró máxima actividad a pH 8,5 y 40°C, usando BAPNA como sustrato. La actividad de la enzima se afectó de manera efectiva por el metil fenil sulfonil fluoruro (PMSF) (98%) y por Na-p-tosil-L-lisina

clorometil cetona (TLCK) (99%). Además la actividad de la enzima fue afectada por los siguientes iones en orden decreciente: Hg^{2+} , Fe^{2+} , Cu^{2+} , Li^{+} , Mg^{2+} , K^{+} , Mn^{2+} , Ca^{2+} . Por otro lado la actividad de la enzima disminuyó de manera progresiva a medida que aumentó la concentración de NaCl de 0 a 30%. Los valores de K_m y k_{cat} fueron de $0,085 \pm 1,45$ mM y $1,76 \pm 0,12s^{-1}$, respectivamente. Según los resultados obtenidos se sugiere que la enzima presenta potencial de uso biotecnológico.

TRIPSINA DO HEPATOPÂNCREAS DE LULA GIGANTE (*Dosidicus gigas*): PURIFICAÇÃO E CARACTERIZAÇÃO

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RESUMO

Foi possível purificar tripsina do hepatopâncreas de *Dosidicus gigas* por fracionamento com sulfato de amônio (30-70% de saturação), cromatografia de filtração em gel, afinidade e troca iônica. O peso molecular da tripsina foi de 23.5 kDa, segundo a eletroforese SDS-PAGE e mostrou uma só faixa no zimograma. A enzima mostrou máxima atividade em pH 8.5 e 40°C, usando BApNA como substrato. A atividade da enzima foi afetada de maneira efetiva por metil-fenil-sulfonil fluoreto (PMSF) (98%) e por Na-p-tosil-L-lisina

clorometil cetona (TLCK) (99%). Também a atividade da enzima foi afetada pelos seguintes íons em ordem decrescente: Hg^{2+} , Fe^{2+} , Cu^{2+} , Li^{+} , Mg^{2+} , K^{+} , Mn^{2+} , Ca^{2+} . Por outro lado, a atividade da enzima diminuiu de maneira progressiva na medida em que aumentou a concentração de NaCl de 0 a 30%. Os valores de K_m e k_{cat} foram de 0.085 ± 1.45 mM e $1.76 \pm 0.12 s^{-1}$, respectivamente. Segundo os resultados obtidos, sugere-se que a enzima apresenta potencial de uso biotecnológico.

California (27°55'N, 110°54'O). Immediately after capture, specimens were stored in an ice bed system and transported to the seafood laboratory, Universidad de Sonora in Hermosillo, Mexico. The time from capture to the dissection process was under 12h.

Preparation of crude enzyme extract

At the laboratory, thawed jumbo squid specimens were dissected, their hepatopancreas removed, immediately frozen and kept -80°C until further analysis. Hepatopancreas (100g) were homogenized with 200ml of 50mM Tris-HCl buffer, pH 7.5, and 0.5M NaCl at 20,000×g for 2min, and then centrifuged at 18,000×g for 30min at 2-4°C. After centrifugation, most lipids (upper layer) were removed manually and the supernatant was filtered through gauze to further remove them. Finally, the supernatant (enzyme extract, without

lipids) was frozen and kept at -80°C until further analysis (Heu *et al.*, 1995).

Enzyme purification

Crude extract was fractionated with ammonium sulfate at 30 and 70% saturation, according to Whitaker (1994). Each time sample was centrifuged at 20,000×g for 20min at 4°C. The precipitate was dissolved in 20ml of 50mM Tris-HCl buffer, pH 7.5 (buffer A), loaded into a 1.6 x 120cm G-75 Sephadex gel filtration chromatography column (Amersham Pharmacia Biotech), eluted at 0.3ml/min and equilibrated with buffer A, collecting 3ml fractions. Then, fractions with trypsin-like activity were pooled and loaded into a 1.1 x 15cm Benzamidine-Sepharose 4 Fast Flow column (Amersham Pharmacia Biotech) equilibrated with the same buffer. The retained fractions were eluted by changing the mobile phase pH from 7.5 to 3.0, using a

50mM Gly-HCl, pH 3.0, buffer (after elution, 200µl Tris-HCl 1M, pH 9.0, were added to each tube). Fractions eluted with trypsin-like activity were combined, dialyzed against 6l of 20mM Tris-HCl, pH 7.5 (buffer B), and pooled for further analysis. For analyses such as optimal pH and temperature (and their respective stabilities), inhibitors and NaCl effect, purified trypsin was used.

Protein concentration determination

Protein concentration in peaks that showed absorbance at 280nm during protein purification was determined according to Bradford (1976) using bovine serum albumin as standard.

SDS-PAGE and zymography

Sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) was carried according to the method of Laemmli

(1970) using 4% stacking gel and 14% separating gel. For protein band analysis, a volume of enzyme extract was mixed with two volumes of sample buffer (125mM Tris-HCl, pH 6.8, 4% SDS, 20% glycerol, 10% β-mercaptoethanol, 0.004% bromophenol blue), heated at 95°C for 4min, cooled immediately and loaded into the gel. After electrophoresis, gels were stained with 0.25% Coomassie brilliant blue R-250 (CBB) in 40% methanol, 10% acetic acid or silver staining, depending on the purification process (protein concentration). Bovine serum albumin (66kDa), ovalbumin (45kDa), glyceraldehyde 3-phosphate dehydrogenase (36kDa), carbonic anhydrase (29kDa), trypsinogen (24kDa), trypsin inhibitor (20kDa) and lactalbumin (14.2kDa) were used as molecular weight markers.

Zymograms were performed according to the procedure of Laemmli (1970) except that samples were not heated and

no reducing agents were added. After electrophoresis, the gel was soaked for 30min in 1.25% casein in 50mM Tris-HCl, pH 8.0, at 4°C. Then, the gel was immersed in the same solution at 37°C for 60min, soaked in trichloroacetic acid for 30min to stop the reaction, washed in distilled water, fixed and stained with 0.05% Coomassie Blue solution, and finally destained with 40% methanol and 10% acetic acid. Native-PAGE was performed according to the procedure of Laemmli (1970) except that samples were not heated and no SDS and reducing agents were added.

Amidase activity

Trypsin amidase activity was evaluated according to Erlanger *et al.* (1961) using 10µl of enzyme extract combined with 990µl N-benzoyl-DL-arginine p-nitroanilide (BAPNA) solution (1mM BAPNA dissolved in 50mM Tris-HCl, pH 7.5 buffer) at 25°C. Production of p-nitroaniline was measured by monitoring the increment in absorbance at 410nm (A_{410}) every 30s for 10min. BAPNA hydrolysis units (U) were calculated as $U = A_{410} / \text{min} \times 1000 \times 1/8,800 \times \text{mg enzyme}$, where 8,800 $\text{M} \cdot \text{cm}^{-1}$ is the p-nitroaniline molar extinction coefficient at 410 nm.

Trypsin esterase activity

Esterase activity was evaluated according to Hummel (1959) using 1mM of N- α -p-tosyl-L-arginine methyl ester hydrochloride (TAME; 50mM Tris-HCl, pH 8.0 buffer) as substrate. Briefly, 10µl of enzyme solution was mixed with 990µl TAME at 25°C. Production of p-tosyl-arginine was measured by monitoring the increment in absorbance at 247nm (A_{247}) every 30sec for 10min. TAME units (U) were calculated as $U = A_{247} / \text{min} \times 1000 \times 1/540 \times \text{mg enzyme}$, where 540 $\text{M} \cdot \text{cm}^{-1}$ is the p-tosyl-arginine molar extinction coefficient at 247nm.

Optimum pH and temperature

The effect of pH on trypsin activity was evaluated using a

universal buffer from pH 4 to 11 at 25°C for 15min (Stauffer, 1989). In order to study the effect of temperature on enzyme activity, the extract was incubated at 20, 25, 30, 35, 40, 50, 55, 60, 65 and 70°C for 15min in 50mM Tris-HCl buffer under optimal pH.

pH and thermal stability

The effect of pH and temperature on enzyme stability was evaluated by measuring the residual activity after incubation at various pH (from 4 to 11) for 60min at 25°C using universal buffers (Stauffer, 1989). Enzyme solution temperature stability was evaluated by incubation at various temperatures (25, 35, 45, 55 and 65°C) for 60min.

Effect of inhibitors

In order to elucidate the effect of inhibitors on trypsin activity, a series of inhibitors were evaluated at different concentrations. Phenylmethylsulfonyl fluoride (PMSF; 100mM in 2-propanol), N- α -Tosyl-L-lysine chloromethyl ketone (TLCK; 10mM in dimethyl sulfoxide, DMSO), benzamidine (10mM in DMSO), soybean trypsin inhibitor ($1\text{g} \cdot \text{l}^{-1}$ in distilled water), N-tosyl-L-phenylalanine chloromethyl ketone (TPCK; 5mM in methanol), ethylenediamine tetraacetic acid (EDTA; 10mM in distilled water) and pepstatin A (1mM in DMSO) were used for the analysis. A mixture of 50µl inhibitor solution and 50µl enzyme extract was incubated for 60min at 25°C; then 950µl of substrate solution (1mM BAPNA in 50mM Tris-HCl buffer, pH 7.5) was added and

residual activity was measured. Control tests were performed in absence of inhibitors.

Effect of metal ions

Samples (30µl) of the purified enzyme were pre-incubated with 70µl of various metal ion solutions (KCl, LiCl, CaCl₂, MnCl₂, MgCl₂, CuSO₄, FeSO₄ and HgSO₄ at 5mM) for 60min at 25°C. Thereafter, 900µl of substrate solution (1mM BAPNA in 50mM Tris-HCl buffer, pH 8.0) was added and the residual activity measured (Souza *et al.*, 2007; Ktari *et al.*, 2012). Control tests were performed in absence of metal ions.

Effect of NaCl on enzyme activity

Trypsin activity was assayed in the presence of NaCl at varying final concentrations (0, 5, 10, 15, 20, 25 and 30% w/v). Residual activity was determined at 25°C at pH 7.5 using 1mM TAME as a substrate. The activity of the enzyme in the absence of NaCl was taken as a control.

Kinetic parameters

The Michaelis-Menten constant (K_m) and catalysis constant (k_{cat}) were evaluated. The initial velocity of the enzymatic reaction was evaluated at 25°C by varying BAPNA substrate concentration (0.01, 0.05, 0.1, 0.25, 0.5, 0.75 and 1.0mM). K_m and V_{max} were calculated from Lineweaver-Burk plots (Lineweaver and Burk, 1934). The value of the turnover number or k_{cat} was calculated from $k_{cat} \times V_{max} / [E]$, where [E] is the active enzyme concentration (Copeland, 2000).

Results

Trypsin purification

Results on purification steps of trypsin from jumbo squid (*Dosidicus gigas*) hepatopancreas are summarized in Table I. As observed, trypsin was purified 59.77 fold, achieving a 63.66% yield. Gel filtration chromatography (G-75 Sephadex) separated the high molecular weight protein group, with low trypsin specific activity, from the low molecular weight protein group showing higher trypsin specific activity. Fractions with high trypsin activity were combined for affinity chromatography (Benzamidine-Sepharose). This last chromatography detected protein in fractions 4 to 10, but with no trypsin specific activity (Figure 1a); however, fractions 40 to 48 showed high trypsin specific activity and were combined for ionic exchange chromatography (DEAE-Sepharose) (Figure 1b). Washing (equilibration) buffer conditions (fractions 1 to 77) showed scarce protein (fractions 4 to 11) with no enzymatic activity detected. The NaCl linear gradient, applied after fraction 73, eluted most of the protein with trypsin activity (fractions 78 to 81).

SDS-PAGE and zymography

Purified trypsin showed a single band on both SDS and zymography (Figure 2a, b) suggesting their high purity and showing only one dominant isoform of trypsin in *Dosidicus gigas* hepatopancreas. Trypsins have been reported to have molecular masses between 20 and 30kDa in mammals and aquatic species. The purified trypsin enzyme from *Dosidicus gigas*

TABLE I
A SUMMARY OF PURIFICATION OF TRYPSIN FROM
THE HEPATOPANCREAS OF *Dosidicus gigas*

Purification step	Total protein (mg)	Total activity (U)	Specific activity (U/mg)	Purification fold	Recovery (%)
Crude extract	7265.35	6102.89	0.84	1	100
Ammonium sulfate	1268.22	5871.86	4.63	5.51	96.21
Gel Filtration	563.14	4775.43	8.48	10.10	81.33
Affinity	145.21	2629.75	18.11	21.56	55.07
Ionic exchange	33.34	1674.00	50.21	59.77	63.66

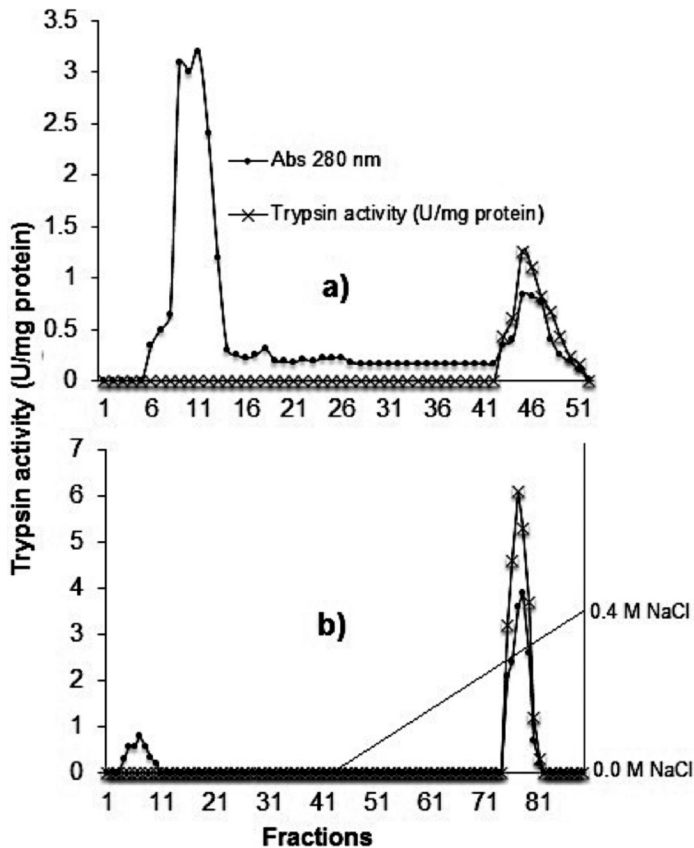


Figure 1. Chromatographic purification of trypsin from hepatopancreas of *Dosidicus gigas*. a: Affinity chromatography. b: DEAE-Sepharose chromatography. Trypsin activity in BAPNA.

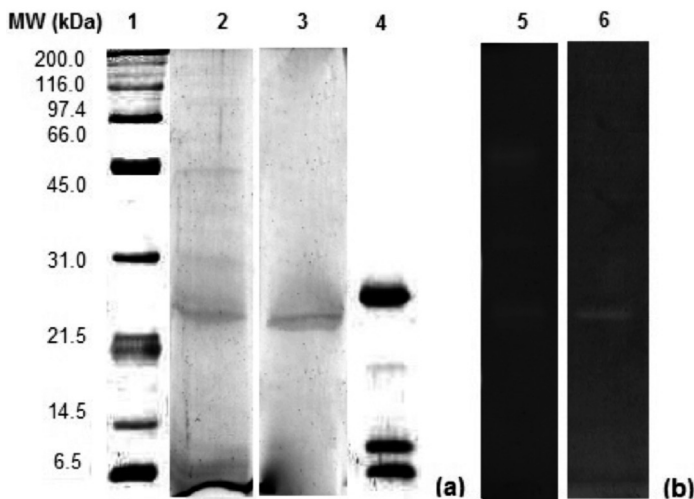


Figure 2. SDS-PAGE (a) and Native-PAGE (b) zymography of purified trypsin from hepatopancreas of *Dosidicus gigas*. Lane 1: molecular weight markers, lane 2: crude extract, lane 3: DEAE-Sepharose fraction, lane 4: bovine trypsin, lane 5: crude extract, lane 6: DEAE-Sepharose fraction.

showed a molecular weight of 23.5kDa, similar to several fish trypsin such as rainbow trout (25.7kDa; Kristjasson, 1991), common carp (28kDa; Cao

et al., 2000), Monterey sardine (*Sardinops sagax caerulea*; 25kDa; Castillo-Yañez *et al.*, 2005), Atlantic bonito (*Sarda sarda*; 29kDa; Klomklao *et al.*,

2007), walleye pollock (24kDa; Kishimura *et al.*, 2008), sardine (*Sardinella aurita*; 28.8kDa; Khaled *et al.*, 2011) and zebra blenny (*Salaria basilisca*; 27kDa; Ktari *et al.*, 2012).

Optimum pH and pH stability

Optimum pH for *Dosidicus gigas* trypsin activity was 8.5 (Figure 3), although high activity was observed throughout the pH range of 6.5 to 11. The optimum pH falls in the range for alkaline aquatic species digestive proteases such as Monterey sardine (*Sardinops sagax caerulea*; pH 8.0; Castillo-Yañez *et al.*, 2005), mandarin fish (*Siniperca chuatsi*; pH 8.5; Lu *et al.*, 2008), silver mojarra (*Diapterus rhombeus*; pH 8.5; Silva *et al.*, 2011), hybrid catfish (*Clarias macrocephalus* × *Clarias gariepinus*; pH 8.0; Klomklao *et al.*, 2011). A considerable loss of activity, although still some of it remained, was observed from pH from 5.5 to 4.0, with 36 to 13% of activity. Trypsin stability at different pH remained remarkable high over a broad pH range, from 7.0 to 11.0, maintaining more than 80% of activity; however, the enzyme was unstable at lower pH (4.5 and 4.0). Similar results were found by (Klomklao *et al.*, 2006; Khishimura *et al.*, 2008; Lu *et al.*, 2008; Jellouli *et al.*, 2009) for tongol tuna (*Thunnus tonggol*), walleye pollock (*Theragra chalcogramma*), man-

darin fish (*Siniperca chuatsi*), and grey triggerfish (*Balistes capriscus*), respectively.

Optimum temperature and thermal stability

The optimum temperature for *Dosidicus gigas* trypsin activity under experimental conditions was 40°C (Figure 4). The same result was obtained for anionic trypsin from grey triggerfish (*Balistes capriscus*; Jellouli *et al.*, 2009) and mandarin fish (*Siniperca chuatsi*; Lu *et al.*, 2008) while higher results were found for cod (*Gadus ogac*; 55°C; Simpson and Haard, 1987), anchovy (*Engraulis encrasicolus*; Martínez *et al.*, 1988), silver mojarra (*Diapterus rhombeus*; 55°C; Silva *et al.*, 2011), hybrid catfish (*Clarias macrocephalus* × *Clarias gariepinus*; 60°C; Klomklao *et al.*, 2011) and zebra blenny (*Salaria basilisca*; 70°C; Ktari *et al.*, 2012). The difference in optimal temperatures might be related to the aquatic species inhabiting environment.

The thermal stability profile (Figure 5) showed that *Dosidicus gigas* trypsin is highly stable at 25 and 35°C, maintaining more than 80% of its activity; however, its activity decreases sharply at temperatures above 55°C. The enzyme was inactivated after 15min of incubation at 65°C. The thermostability profile shown by this enzyme is similar to that of other fish trypsins, like sardine (*Sardinops sagax*

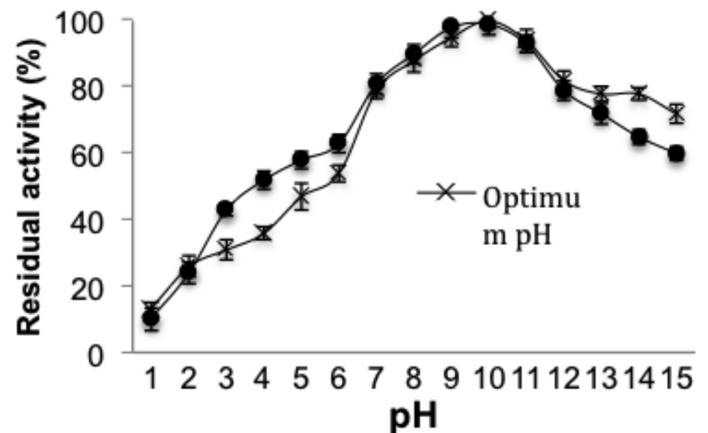


Figure 3. Optimum pH and pH stability of purified trypsin from viscera of *Dosidicus gigas*.

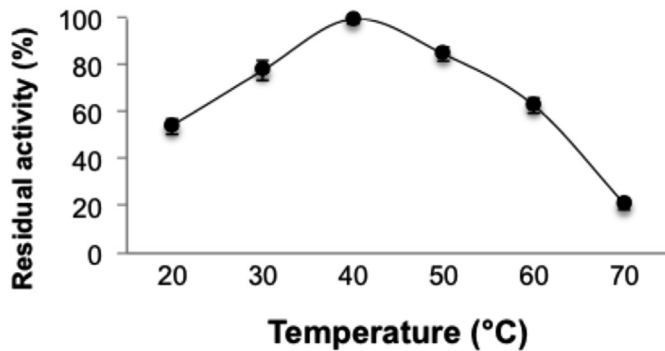


Figure 4. Optimum temperature of purified trypsin from hepatopancreas of *Dosidicus gigas*.

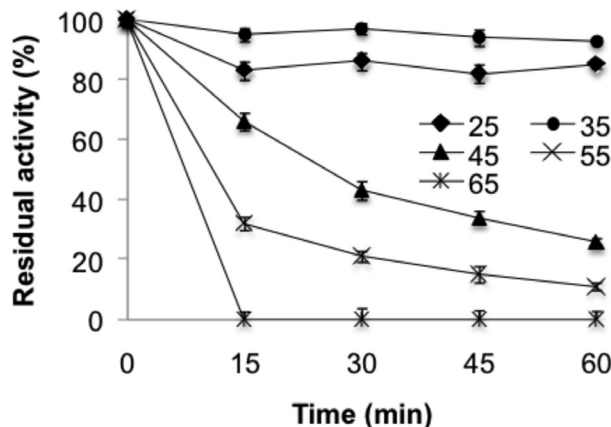


Figure 5. Thermal stability of purified trypsin from hepatopancreas of *Dosidicus gigas*.

caerulea; Castillo-Yañez *et al.*, 2055), bogue (*Boops boops*; Barkia *et al.*, 2010) and zebra blenny (*Salaria basilisca*; Ktari *et al.*, 2012).

Effect of inhibitors and metal ions

Proteases can be classified by their sensibility to various inhibitors (North, 1982). The effect of several proteinase inhibitors on *Dosidicus gigas* trypsin activity is summarized in Table II. The enzyme was 98% inhibited by PMSF, a serine-protease inhibitor. Trypsin activity was strongly inhibited by TLCK (83%) and benzamidine (85%), both specific inhibitors of trypsin. EDTA (a metallo protease inactivator), TPCK (a chymotrypsin inhibitor) and pepstatin A (an aspartic proteases inhibitor) did not affect trypsin activity (<10% inhibition). These results con-

firm that the single band detected by SDS and zymogram corresponds to trypsin. Similar results were observed for trypsin from Monterrey sardine (Castillo-Yañez *et al.*, 2005), Pacific cod (*Gadus macrocephalus*; Fuchise *et al.*, 2009) and brownstripe red snapper (*Lutjanus vitta*; Khantaphant and Benjakul, 2010).

The effect of metal ions over enzyme activity is also shown in Table II. It can be observed that Ca^{2+} ion did not have an effect over proteolytic activity; however, the enzyme was slightly inactivated by Mn^{2+} , K^+ , Mg^{2+} , Li^+ and Cu^{2+} ions, down to 94, 91, 82, 78 and 68% residual activity, respectively. On the other hand, Fe^{2+} and Hg^{2+} ions greatly affected trypsin activity, showing only 29 and 23% residual activity, respectively. Similar results were observed in a trypsin from *Sardinella aurita* (Khaled

TABLE II
EFFECT OF PROTEASE INHIBITORS AND METAL IONS ON THE ACTIVITY OF TRYPSIN FROM *Dosidicus gigas*

Inhibitor/Ion	Concentration	Residual activity (%)
Control (No inhibitor/Ion)		100
PMSF	100 mM	2 ±0.23
TLCK	10 mM	0.8 ±0.16
Benzamidine	10 mM	5 ±0.37
TPCK	5 mM	95 ±0.82
EDTA	2 mM	96 ±1.03
Pepstatin A	0.1 mM	95 ±0.62
CaCl_2	5 mM	93 ±1.22
MnCl_2	5 mM	94 ±0.78
KCl	5 mM	91 ±0.51
MgCl_2	5 mM	82 ±0.45
LiCl	5 mM	78 ±1.32
CuSO_4	5 mM	68 ±1.83
FeSO_4	5 mM	29 ±0.40
HgSO_4	5 mM	23 ±0.82

et al., 2011) and bogue (*Boops boops*; Barkia *et al.*, 2010). It has been reported that Hg^{2+} ions bind to -SH groups on the enzymes, inhibiting their action (Klee, 1988).

Effect of NaCl on enzyme activity

The effect of NaCl on the activity of *Dosidicus gigas* trypsin showed a continuous decreased in activity with increasing NaCl concentration. However, it remained very active even at high salt concentrations, showing 87, 72, 61, 53, 44 and 32% residual activity at 5, 10, 15, 20, 25 and 30% of NaCl, respectively.

Kinetic studies

The kinetic constants, K_m and k_{cat} of purified *Dosidicus gigas* trypsin, hydrolyzing BAPNA at 25°C, were determined using a Lineweaver-Burk plot. The values of K_m and k_{cat} presented by the purified trypsin were 0.085mM and 1.76s⁻¹, respectively. K_m was similar to that reported for *B. caprisicus* (0.07mM; Jellouli *et al.*, 2009), sardinelle (*Sardinella aurita*; 0.125mM; Khaled *et al.*, 2009), but lower than those from snapper (*P. macracanthus*; Hau and Benjakul, 2006) and brownstripe red snapper (*Lutjanus vitta*; Khantaphant and Benjakul, 2010). The k_{cat} value was similar to those found in trypsin from anchoveta, of 1.55s⁻¹ and snapper, of

1.06s⁻¹ (*E. japonica*; Heu *et al.*, 1995 and *P. macracanthus*; Hau and Benjakul, 2006).

Discussion

Based on the behavior observed during DEAE-Sephrose separation, it is concluded that the retained protein, with trypsin activity, is of anionic nature. It has been observed that trypsin from fish viscera have high ratios of acidic to basic amino acids. This characteristic is quite different from mammalian trypsin, which are of basic nature (Heu *et al.*, 1995). Anionic trypsin are common in fish such as salmon (*Oncorhynchus keta*), sardine (*Sardinops melanosticta*), anchovy (*Engraulis encrasicolus*), sardine (*Sardinops sagax caerulea*) and pez diablo (*Pterygoplichthys disjunctivus*) (Martínez *et al.*, 1988; Sekisaki *et al.*, 2000; Castillo-Yañez *et al.*, 2005; Villalba-Villalba *et al.*, 2013). The differences in trypsin molecular mass may be due to genetic variation among species, but the possibility that these differences are caused by autolytic degradation should not be excluded (Lu *et al.*, 2008).

Trypsins generally belong to the alkaline proteinase group (Simpson and Haard, 1987); thus, under acid conditions the charge distribution and conformation were changed and the enzyme could not bind to substrate properly (Benjakul and Morrissy, 1997). The stability

of enzymes at a particular pH may be related to its net charge at that pH; at low pH, the increment of positive charges on the enzyme destabilizes them, affecting its tertiary structure (Castillo-Yañez *et al.*, 2006). *Dosidicus gigas* trypsin showed a high activity and stability at high alkaline pH conditions, as well as in slightly acid ones, which makes it a potential candidate for application in food processing operations or for its applications in detergents and food industry.

That the trypsin showed low thermostability could be related to a high proportion of charged residues and fewer polar hydrogen-bond forming residues as suggested by Benjakul and Morrissey, 1997. High activity of trypsin from *Dosidicus gigas* at low temperatures may be interesting for many biotechnological and food protein processing applications. Besides, this low thermostability can be advantageous, as these enzymes can be easily inactivated by using less heat treatment (Jiang *et al.*, 2010).

The inhibitors employed are highly specific, possessing at their surface one or more peptide bonds made with the α -carboxylic side of arginine known as reactive sites. The reactive site specifically interacts with the active site of the cognate enzyme. Both natural and synthetic inhibitors inhibit trypsin. The extent of the inhibition (K_i = molar) for trypsin goes from millimolar to femtomolar, indicating that quite different affinities between them are possible (Zollner, 1993). In the present study, the pattern of inhibition observed in inhibitors is characteristic of other trypsins, supporting the finding that the isolated enzymes is, indeed, a trypsin.

Loss of enzyme activity due to the denaturation of trypsin caused by the 'salting out' effect (Khaled *et al.*, 2011) could be observed. Similar results were found in trypsin from bogue (*Boops boops*; Klee, 1998). Results show that trypsin from *Dosidicus gigas* may have potential to be used to accelerate hydrolysis of

proteins under hypersaline conditions, such as in fermented products like fish sauce.

The catalytic efficiency (k_{cat}/K_m) value of *Dosidicus gigas* trypsin (20.71 s \cdot mM $^{-1}$), compared with that of bovine trypsin (3.1 s \cdot mM $^{-1}$; Asgeirsson *et al.*, 1989) revealed a higher catalytic efficiency (6.68 folds at 25°C), which means that *Dosidicus gigas* trypsin is more prompt to attach to and hydrolyze the substrate than the bovine type.

Trypsins shows species-specific characteristics, since there are significant differences in feeding habits, ingested food composition and differences in the protein digestion process, according to the requirements of the species. These differences are adaptive responses to different life styles, environments and mostly, different survival abilities among species. However, trypsins share some general characteristics, mainly related to their catalytic features (Muhlia-Almazán *et al.*, 2008).

Conclusions

This paper describes the characteristics of a serine protease obtained from *Dosidicus gigas* hepatopancreas. Based on the results, it can be concluded that the isolated and purified enzyme was trypsin, showing high activity at pH 6.5-11.0, 25-40°C and 30% salt concentration. Therefore, the hepatopancreas of *Dosidicus gigas* can be an important source of trypsin, whose pH stability, activity under high salt concentration and thermal behaviour make it a potential agent to be used in biotechnological processes.

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