

Isolation of a thermostable trypsin inhibitor with exploitable potential

Yongxiang Hong · Xixi Cai · Biao Shao ·
Jing Hong · Shaoyun Wang · Pingfan Rao

Received: 28 October 2012/Revised: 20 April 2013/Accepted: 28 April 2013/Published online: 15 May 2013
© Springer-Verlag Berlin Heidelberg 2013

Abstract A novel trypsin inhibitor with considerable thermal and pH stability, designated *Glytine*, was isolated from seeds of the Chinese black soybean *Glycine max* (L.) Merr. The purification procedure involved ammonium sulfate precipitation, ion-exchange chromatography on CM-Sephadex C-50, gel filtration chromatography on Sephacryl S-200HR, and gel filtration chromatography on POROS HS-20. The 20 N-terminal amino acid sequences were determined to be DEYSKPCCDLCMCTRRCPPQ, demonstrating close homology with the sequences of leguminous trypsin inhibitors. The molecular mass and isoelectric point of the inhibitor were estimated by SDS-PAGE and isoelectric focusing to be 19.9 kDa and 6.2, respectively. Trypsin could be completely inhibited by *Glytine* when the weight ratio was 1.5. The inhibitory activity of *Glytine* was unaffected by exposure to temperatures up to 100 °C, or within the pH range 2–12. Besides trypsin–chymotrypsin inhibition activity, *Glytine* demonstrated other biological activities including antiproliferative activity against tumor cells including human liver hepatoma cells Bel-7402 and neuroblastoma cells SHSY5Y. In addition, the inhibitor showed antifungal activity against *Pythium aphanidermatum*, *Fusarium oxysporum*, *Alternaria alternata* (Fr.) Keiss, *Fusarium solani*, and *Botrytis cinerea*. This study extended research on leguminous

trypsin–chymotrypsin inhibitor and suggested exploitable potential.

Keywords Trypsin inhibitor · Thermostable · *Glycine max* (L.) Merr · Antifungal

Introduction

Leguminous plants elaborate a range of proteins with diverse bioactivities. Protease inhibitor is listed as one type of these bioactive proteins having been researched by a number of investigators. Besides inhibiting the activity of protease, protein inhibitors of serine including trypsin and chymotrypsin concurrently demonstrated additional biological activities. For example, HIV protease inhibitors and SARS coronavirus proteinase inhibitors may be used to fight against HIV and SARS virus, respectively [1]. Plant protease inhibitors may be involved in the regulation of programmed cell death in plants. They show insecticidal and antifungal activities, but can also cause inhalant and food allergies [2, 3]. Epidemiological evidence suggests that a diet rich in legumes reduces the incidence of cancer and that this may be due to the presence of protease inhibitors [4]. Both in vivo antitumor and in vitro antiproliferative activities of trypsin inhibitors have been reported [5].

There are several types of plant trypsin inhibitors. Kunitz-type trypsin inhibitors have a molecular mass of about 20 kDa, a low cysteine content and a single reactive site, while Bowman–Birk trypsin inhibitors are approximately 8 kDa in size and possess a high cysteine content and two reactive sites [6, 7]. The conformation of Kunitz-type trypsin inhibitors is mainly β sheet with a small amount of regular sheet. The insecticidal activity of Kunitz-type

Yongxiang Hong and Xixi Cai contributed equally to this work.

Y. Hong · X. Cai · J. Hong · S. Wang (✉) · P. Rao
College of Bioscience and Biotechnology, Fuzhou University,
Fuzhou 350108, People's Republic of China
e-mail: shywang@fzu.edu.cn

B. Shao
Nantong Products Quality Supervision and Inspection Institute,
Nantong 226011, People's Republic of China

trypsin inhibitors has been demonstrated using transgenic plants [8]. Kunitz-type trypsin inhibitors also inhibit other enzymes, such as chymotrypsin, α -amylase, and human plasmin, and block the conversion of prothrombin to thrombin [3]. The Yellow soybean (*Glycine max* (L.) Merr.) produced both Bowman–Birk and Kunitz-type trypsin inhibitors [9]. The formation, degradation, and gene expression of Kunitz-type trypsin inhibitor in the soy bean have also been reported [3]. The seeds of bitter gourds, sponge gourds, wax gourds, and *M. cochinchinensis* produce squash-type trypsin inhibitors with a molecular mass of about 3 kDa [10].

Trypsin and chymotrypsin inhibitors have been regarded as a class of antifungal proteins that have potent activity against plant pathogens [10]. Antifungal proteins have captured the attention of a number of researchers on account of their important economic potential in protecting crops from invading fungi. Antifungal proteins may also contribute to the defense of a plant against predators such as insects [11] as well as pathogens such as fungi [12, 13]. It is well recognized that there is a wide diversity of antifungal proteins produced by leguminous plants. Some of these proteins and peptides have been classified into different groups based on their structure and/or functions [14]. Combinations of antifungal proteins may be found in a single species of bean; for example, in addition to a protease inhibitor, a lysozyme and a nonspecific lipid transfer protein were isolated from Mung bean (*Vigna radiata*), and all have fungal inhibition activity [12, 13, 15].

The Chinese black soybean is a special cultivar of the Yellow soybean (*Glycine max* (L.) Merr). Its usage in traditional Chinese medicine is different from the Yellow soybean. However, Chinese black soybean has not been studied as extensively as the Yellow soybean, and there is therefore a paucity of information about its protein constituents. We report here for the first time the extraction from it, and purification, of a trypsin inhibitor with both antifungal activity and antiproliferative activity against tumor cells.

Materials and methods

Materials

Black soybean (*Glycine max* (L.) Merr) seeds were purchased from a local supermarket. Fetal calf serum (FCS) was purchased from Hyclone Co., USA. The human liver hepatoma cells of Bel-7402 and the neuroblastoma cells of SHSY5Y were obtained from Shanghai Cell Institute, China Science Academy, China. Trypsin and chymotrypsin from bovine pancreas were purchased as white crystalline powder from Shanghai Chemical Co. (Shanghai, China);

their enzymatic activities were 2,400 and 800 μ /mg, respectively.

CM-Sephadex C-50, Sephacryl S-200HR, and POROS HS-20 were purchased from Amersham Biosciences (Sweden), Bio-Rad (USA), and TOSOH Co. (Japan), respectively. Standard proteins marker were purchased from Gibco-BRL (Life Tech., USA).

Sample preparation

Hundred grams of Black soybeans seeds were soaked in distilled water for 12 h and homogenized in 0.2 M sodium acetate buffer, pH 5.4. The homogenate was centrifuged at 12,000 rpm for 20 min at 4 °C. The supernatant was collected for further study.

Ammonium sulfate precipitation

The supernatant was first brought to 20 % saturation with solid ammonium sulfate. The solution was centrifuged at 12,000 rpm for 20 min, and the resulting supernatant was then adjusted to 80 % saturation by the addition of ammonium sulfate. After centrifugation (12,000 rpm, 20 min), the precipitate was collected and dissolved in 300 mL of 0.02 M sodium acetate buffer (pH 5.4). The solution of ammonium sulfate precipitate was dialyzed against 0.02 M sodium acetate buffer (pH 5.4) overnight at 4 °C with three changes.

Cation-exchange chromatography

The dialyzed solution was applied to a CM-Sephadex C-50 column (2.5 cm \times 30 cm) equilibrated previously with the 0.02 M sodium acetate buffer (pH 5.4). The column was washed with the binding buffer and eluted with a 0–0.5 M linear gradient of NaCl. The flow rate was 0.5 mL/min, 10 min/tube. The absorbance of the elution was monitored at 280 nm. The trypsin inhibitor activity of each fraction was determined. The unabsorbed peak (fraction C₁) exhibited trypsin inhibitor activity.

Gel filtration

The unabsorbed fractions from the CM-Sephadex C-50 column were pooled, dialyzed against 0.01 M Tris–HCl buffer (pH 7.2) at 4 °C for 24 h with several changes and concentrated using a vacuum concentration machine (Shanghai Machinery Co., China) at low temperature for 2 h, and then applied to the Sephacryl S-200HR column (1 cm \times 100 cm) previously equilibrated with the starting buffer. The flow rate was 0.3 mL/min, and the eluate was monitored at 280 nm. The trypsin inhibitor activity of each fraction was determined.

Anion-exchange chromatography

The pooled active solution was dialyzed against 0.02 M acetate buffer (pH 5.0) and loaded onto a POROS HS-20 column (0.75 cm × 7.5 cm). A bound protein was eluted with a linear gradient of 0–1.0 M NaCl for 90 min, and then finally with a stepwise gradient of 1.0 M NaCl for 10 min. Chromatography was carried out on a BioCAD 700E perfusion chromatography workstation from PerSeptive Biosystem Co. (USA) with a flow rate of 1 mL/min. The absorbance of all fractions was monitored at 280 nm, and the trypsin inhibitor activity of each fraction was determined.

Capillary liquid chromatography

The homogeneity of the purified sample was determined on a C18 capillary reversed-phase high-performance liquid chromatography (RP-HPLC) column (15 mm × 1.0 mm) using an analyzer (Applied Biosystems Model ABI-140D, Perkin Elmer Co., MA, USA). The running solvent A contained 90 % water, 10 % acetonitrile, and 0.1 % trifluoroacetic acid, and the solvent B contained 90 % acetonitrile, 10 % water, and 0.1 % trifluoroacetic acid. The gradient eluent concentration was 0–50 % solvent B in 1 h.

Protein assay

The protein concentration was measured by the method of Lowry [16], with bovine serum albumin used as the standard.

SDS-PAGE and IEF

Sodium dodecyl sulfate–polyacrylamide gel electrophoresis (SDS–PAGE) was carried out using a 12.5 % separating gel and 4.0 % concentrating gel, according to the method of Laemmli [17]. Polyacrylamide gel isoelectric focusing (IEF) was conducted in a glass column, the carrier ampholytes (Ampholine, Pharmacia) with pH ranges of 3–10 [18]. Gels were stained by Coomassie blue R-250.

Mass spectrometry

The trypsin inhibitor was analyzed by MALDI-TOF mass spectrometry (Applied Biosystems, Foster City, CA, USA), an instrument that essentially consists of an atmospheric pressure electrospray positive-ion source, attached to a triple-quadrupole mass analyzer.

Measurement of trypsin–chymotrypsin inhibitory activity

Ten portions containing 0, 25, 50, 75, 100, 150, 200, 250, 350, and 450 µg of the inhibitor were incubated at 25 °C together with 25 µg trypsin or chymotrypsin in 100 µL of 0.05 M Tris–HCl buffer (pH 8.0) containing 0.2 M CaCl₂ for 5 min. The reaction was incubated at 25 °C for 15 min, and the reaction was terminated by adding 1 mL of cold 5 % trichloroacetic acid. The reaction mixture was centrifuged for 20 min at 9,000 rpm, and the supernatant was removed for testing. Residual trypsin or chymotrypsin activity was determined by adding 300 µL of 1 % casein substrate at 25 °C. The absorbance of the clear supernatant was determined at 280 nm.

N-terminal amino acid sequence analysis

The N-terminal amino acid sequence of the purified *Glytine* was performed by Edman degradation using a protein sequencer (Applied Biosystems Model 476A, Perkin Elmer Co. MA, USA). Phenylthiohydantoin derivatives were separated and identified by capillary reversed-phase HPLC in a C18 column with an analyzer.

Thermostability assay

Ten portions of 200 µg trypsin inhibitor in 1 mL of glass tubes were preheated at different temperatures (from 30 to 100 °C) for 30 min and cooled down to 4 °C. The residual trypsin inhibition activity was determined according to the method established by Wang and Rao [19].

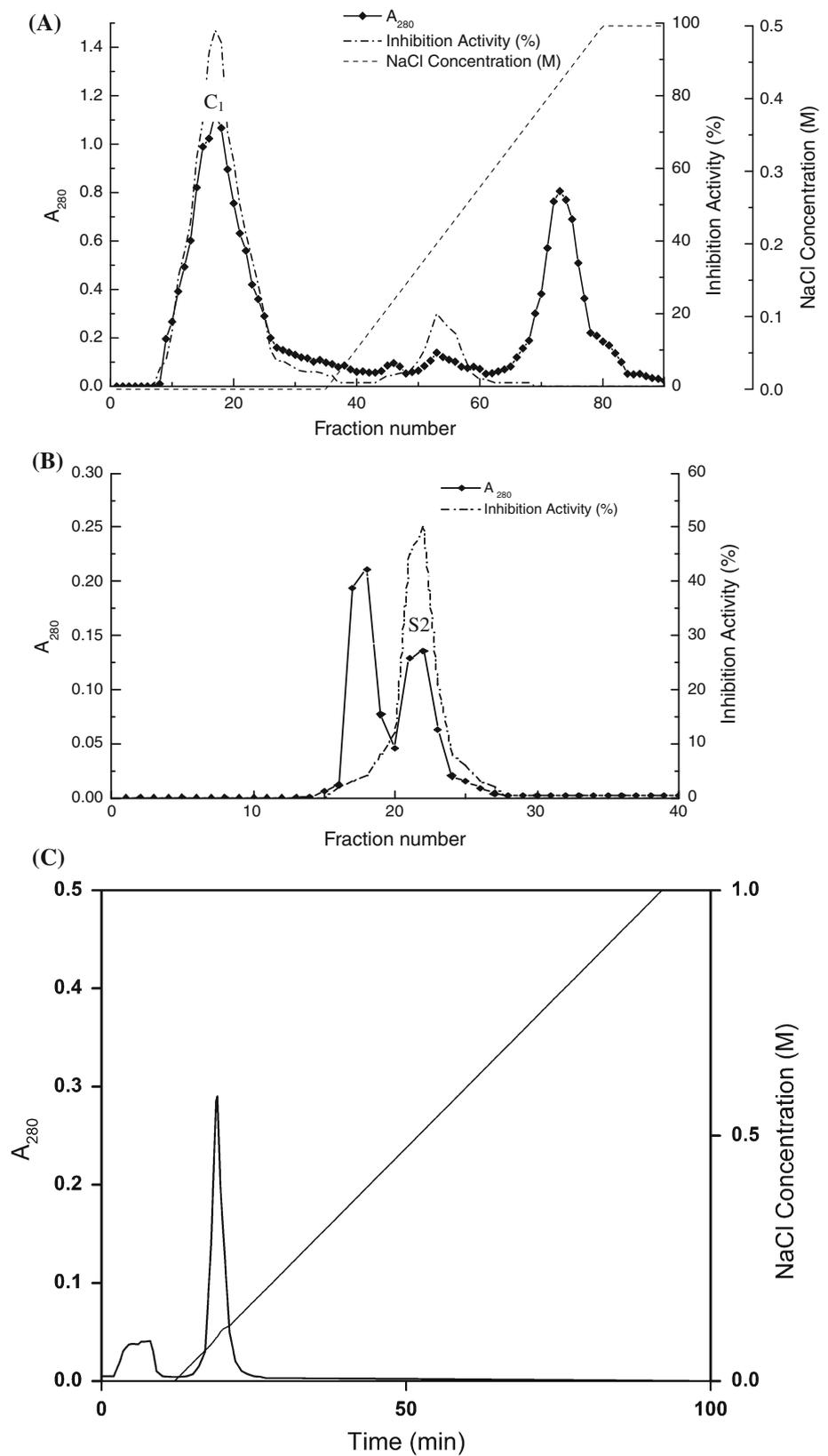
pH stability assay

The purified *Glytine* was incubated in 0.05 M buffers with pH values between 2 and 12 at 25 °C for 30 min. The residual trypsin inhibition activity was determined.

Assay for antifungal activity

The assay for antifungal activity was executed using 100 × 15 mm Petri plates containing 10 mL of potato dextrose agar. Sterile blank paper disks (0.625 cm in diameter) were placed at a distance of about 1 cm away from a central disk of the same size. An aliquot (8 µL containing 0, 50 or 100 µg) of *Glytine* in 0.02 M Tris–HCl buffer (pH 7.2) was introduced to each peripheral disk, respectively. The plates were incubated at 23 °C for 72 h until mycelial growth from the central disk had enveloped the peripheral disks containing the control (Tris–HCl buffer) and had produced crescents indicating inhibition

Fig. 1 Purification of *Glytine*. **a** Cation-exchange chromatography on CM-Sephadex C-50 (2.5 cm × 30 cm). The first elution peak (fraction C₁) was pooled. **b** Fraction C₁ was concentrated and subjected to gel filtration chromatography on Sephacryl S-200HR (1 cm × 100 cm). The second peak (fraction S₂) was collected. **c** Fraction S₂ was concentrated and dialyzed against 0.02 M acetate buffer (pH 5.0) and loaded on to a POROS HS-20 column (0.75 cm × 7.5 cm)



around the disks containing samples with antifungal activity.

Assay for antiproliferative activity

The antiproliferative activity of *Glytine* on tumor cells was carried out by testing its inhibition of the growth of human liver hepatoma cells (Bel-7402) and neuroblastoma cells (SHSY5Y). Whey protein isolate was used as a control to exclude the general effect of protein. The Bel-7402 cells were cultured in RPMI-1640 medium supplemented with 10 % (v/v) FCS. The SHSY5Y cells were cultured in DMEM/F12 (1:1) medium supplemented with 10 % (v/v) fetal bovine serum, in a humidified atmosphere of 5 % CO₂ at 37 °C. The cells (3×10^5 cells/150 μ L/well) were also seeded into a 96-well culture plate, and a series of solutions containing *Glytine* in 150 μ L medium were added. After incubation of the cells at 37 °C in a humidified atmosphere of 5 % CO₂ for 24 h, the cells were then harvested and dyed with 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT). The absorbance of the samples at 590 nm was determined using a microtiter plate (ELISA) reader and was directly correlated to the level of its anti-tumor activity. The inhibitory activity of the *Glytine* was calculated as the percent inhibition compared to a control without the sample.

Statistical analyses

All experiments were conducted in triplicate. All data were presented as means (standard deviations, SDs) of three independent experiments. Statistical analysis was done using Student's *t* test. A value of $P < 0.05$ was considered statistically significant.

Results and discussion

Extraction and purification of *Glytine*

The extract of the Black soybean seeds was firstly treated by ammonium sulfate precipitation (20–80 % saturation) and dialyzed against 0.02 M sodium acetate buffer (pH 5.4). The solution was applied to cation-exchange chromatography on CM-Sephadex C-50 column. The unabsorbed fraction (C₁) exhibiting trypsin inhibitor activity was pooled (Fig. 1a). In the next step, the pooled fraction was subjected to gel filtration chromatography on Sephacryl S-200HR column, and the second peak (fraction S₂; Fig. 1b) with trypsin inhibitory activity was collected and dialyzed against 0.02 M acetate buffer (pH 5.0) and loaded on to a POROS HS-20 column. The trypsin inhibitor was separated from other proteins by eluting with a 0–1.0 M

linear gradient of NaCl (Fig. 1c). The homogeneity of the purified *Glytine* was identified by the C18 capillary reversed-phase chromatography and SDS-PAGE. A single peak and a single band were performed on chromatogram map (Data not shown) and SDS-PAGE (Fig. 2), indicating the high purity of the *Glytine*.

Molecular mass and isoelectric point

The SDS-PAGE pattern is shown in Fig. 2, as a single band on the gel. The molecular weight of the purified *Glytine* was calculated by SDS-PAGE to be 19.9 kDa. The

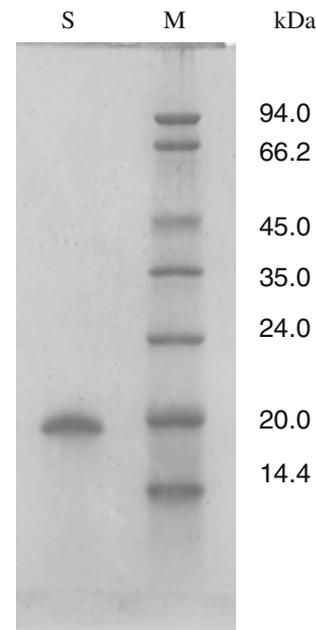
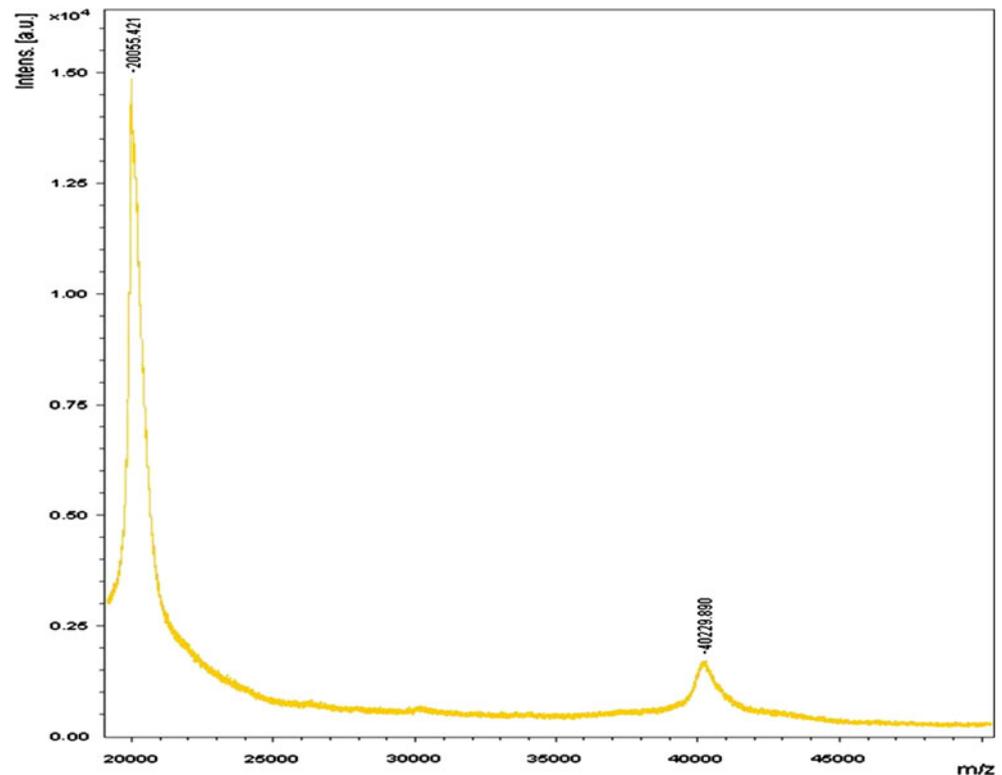


Fig. 2 SDS-PAGE of the purified *Glytine*. From right to left, Lane M was molecular mass standards; lane S was loaded with 8 μ g *Glytine*. The position of *Glytine* band corresponding to a molecular mass of 19.9 kD

Fig. 3 IEF electrophoresis profile of obtained trypsin inhibitor



Fig. 4 MALDI-TOF mass spectrum of obtained trypsin inhibitor



isoelectric point (*pI*) of the purified *Glytine* was estimated to be 6.2 by IEF, suggesting it to be an acidic protein (Fig. 3).

Mass spectrometry

Mass spectrometric analysis on MALDI-TOF mass spectra demonstrated that the molecular mass of the trypsin inhibitor was 20.055 kDa (Fig. 4). This indicates not only its precise molecular mass, but also provides further evidence of the purity of the purified *Glytine*.

N-terminal amino acid sequence

The 20 N-terminal amino acid sequence of the purified trypsin inhibitor *Glytine* was determined to be DEYSKPCCDLCMCTRRCPPQ. This revealed a high correspondence (between 55 and 90 %) to those protease inhibitors and inhibitor precursors from other leguminous plants, according to the results of a BLAST Search (Table 1).

From Table 1, the N-terminal sequence of the *Glytine* was homologous to those of trypsin inhibitors from other

Table 1 Comparison of N-terminal sequence of *Glytine* with trypsin protease inhibitors from other leguminous plants

Source	Isoinhibitor name	Isoinhibitor number	N-terminal sequence	Residue number	Identity (%)
<i>Glycine max</i>		1	DEYSKPCCDLCMCTRRCPPQ	20	100
<i>Glycine soja</i>	Bowman–Birk type proteinase	43	<u>DEYSKPCCDLCMCTR</u> SMPQ	62	90
<i>Glycine max</i>	Bowman–Birk type proteinase	37	<u>DEYSKPCCDLCMCTR</u> SMPQ	56	90
<i>Glycine soja</i>	Bowman–Birk type proteinase	41	<u>DESSKPCCDLCMCT</u> ASMPQ	60	80
<i>Phaseolus microcarpus</i>	Double-headed trypsin inhibitor	51	<u>ESSKPCCDQCACT</u> RSIPPQ	69	80
<i>Phaseolus parvulus</i>	Double-headed trypsin inhibitor	13	<u>ESSEPCCDLCLCT</u> KSIPPQ	31	60
<i>Vigna angularis</i>	Protease inhibitor	8	<u>ESSKPCDECKCT</u> KSEPPQ	26	60
<i>Lens culinaris</i>	Trypsin inhibitor	1	<u>ESSEPCDSCICT</u> KSIPPQ	19	55

The presented N-terminal sequences in Table are from the NCBI database (<http://www.ncbi.nlm.nih.gov>). Residue number 1 and residue number 20 for peanut *Glytine* refer to D and Q being the 1st and 20th amino acid residue in *Glytine*, respectively. Underlined characters, identical amino acids with the purified *Glytine*

leguminous plants. It inhibited the activity of both trypsin and chymotrypsin, with the potency toward trypsin being higher. This result is similar to the Kunitz-type trypsin inhibitors from the roots of Hai Er Shen (Kid Ginseng—*Pseudostellaria heterophylla*) [2] and wild soybean (*Glycine soja*) [20].

Trypsin and chymotrypsin inhibitory activities

Glytine inhibited both trypsin and chymotrypsin activity (Fig. 5). When the weight ratio of inhibitor to chymotrypsin equaled 2.1, 50 % of enzyme activity was inhibited. However, *Glytine*'s capacity to inhibit trypsin activity was much stronger than that to inhibit chymotrypsin activity; when the weight ratio of the *Glytine* to chymotrypsin equaled 2.0, the residual enzyme activity was hardly evident (Fig. 5).

Thermal and pH stability assay

Purified *Glytine* was extremely stable below 100 °C. Incubation at 100 °C for 30 min resulted in almost no loss of activity (Fig. 6). Additionally, it showed excellent pH stability; its trypsin inhibitory activity was unaffected by exposure to pH 2–12 (Fig. 7).

The results show that the thermostability of purified *Glytine* is higher than those of the trypsin inhibitors reported by Lin and Ng [20], Wang et al. [13], and Wang and Rao [19]. The thermostable temperature of almost all of the previously reported plant trypsin inhibitors was lower than 85 °C [19]. The pH stability of *Glytine* is similar to that of the trypsin inhibitor reported by Lin and Ng [20].

Although trypsin inhibitors have been extensively reported, the fact that they differ in both their thermostability and pH stability highlights their diversity, even from the same species. This newly purified trypsin inhibitor, *Glytine*, was both extreme heat-resistant and pH-resistant properties (Figs. 6, 7) when compared with almost all previously reported trypsin inhibitors.

Antifungal activity

The antifungal activity of *Glytine* against fungal species was illustrated in Fig. 8. It showed antifungal activity against *Pythium aphanidermatum* (Fig. 8I), *Fusarium oxysporum* (Fig. 8II), *Alternaria alternata* (Fr.) Keiss (Fig. 8III), *Fusarium solani* (Fig. 8IV), and *Botrytis cinerea* (Fig. 8V). Crescents of inhibition around disks were shown in the position of both Disk (B) and Disk (C) when

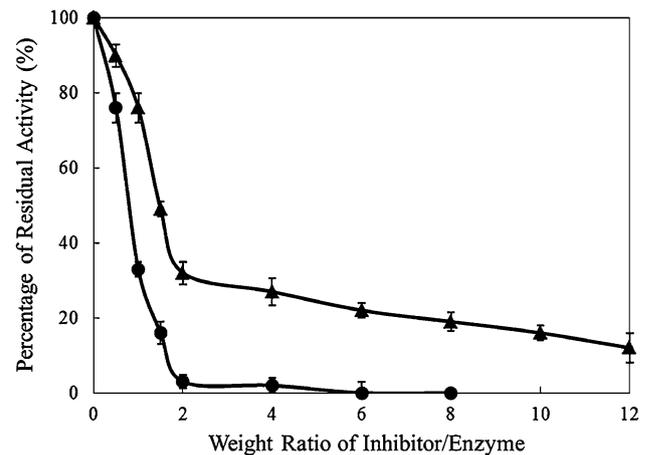


Fig. 5 Inhibitory activity of the trypsin inhibitor *Glytine* on trypsin or chymotrypsin (circle and triangle represent trypsin and chymotrypsin, respectively)

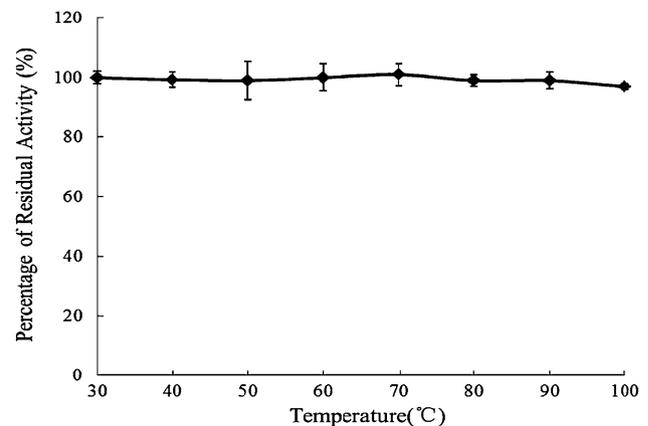


Fig. 6 Thermal stability of the *Glytine*. Results are presented as mean \pm SD ($n = 3$)

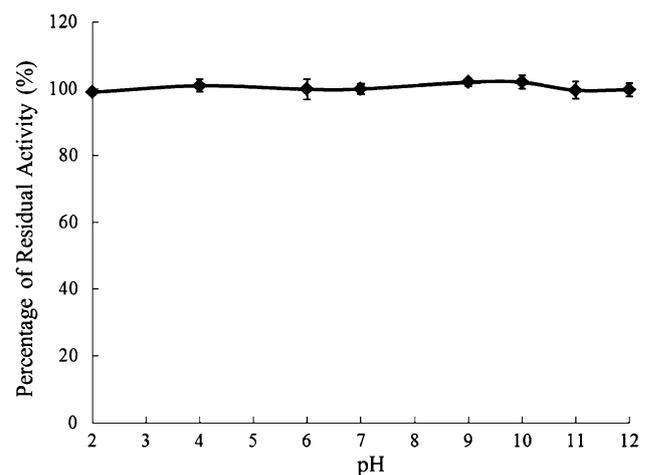


Fig. 7 pH stability of the *Glytine*. Results are presented as mean \pm SD ($n = 3$)

compared with control Disk (A). Moreover, inhibition capacity varied between fungal species; the inhibition of *Botrytis cinerea* (Fig. 8V) was the most visible as compared with Fig. 8I, II, III, and IV. In addition, the IC_{50} of the antifungal activity toward *Pythium aphanidermatum* was calculated to be $9.13 \mu\text{M}$. However, *Glytine* showed little antifungal activity against *Mycosphaerella arachidicola*.

The trypsin inhibitor *Glytine* displays a spectrum of activities. It inhibits mycelial growth in several fungal species, like some of the previously reported antifungal proteins [2, 13, 15, 21, 22]. The existence of trypsin inhibitors with antifungal activity might represent a selective advantage against a wide range of potential pathogens activity. Plant protease inhibitors show antifungal activities; instead, they are just listed as inhalant allergies and also food allergies. It deduces that a combination of antifungal proteins is present in this variety of plants and that they work together to defend against the attacks from invading pathogens such as disease-causing fungi.

Antiproliferative activity on tumor cells

The antiproliferative activities of *Glytine* to human liver hepatoma cells (Bel-7402) and neuroblastoma cells (SHSY5Y) were calculated as percent inhibition compared to a control without the sample. According to the inhibition results (Fig. 9), the IC_{50} value against Bel-7402 was calculated to be almost $40 \mu\text{M}$, while the IC_{50} value against SHSY5Y was near $25 \mu\text{M}$. Whey protein isolate was used as the control to exclude the general effect of protein. Not

Fig. 8 Inhibitory activity of trypsin inhibitor *Glytine*. Plate (1) *Pythium aphanidermatum*, plate (2) *Fusarium oxysporum*, plate (3) *Alternaria alternata* (Fr.) Keiss, plate (4) *Fusarium solani*, and plate (5) *Botrytis cinerea*. Disk (a) 0.02 M Tris–HCl buffer, pH 7.2, as control; disk (b) $50 \mu\text{g}$ *Glytine* and disk (c) $100 \mu\text{g}$ *Glytine*

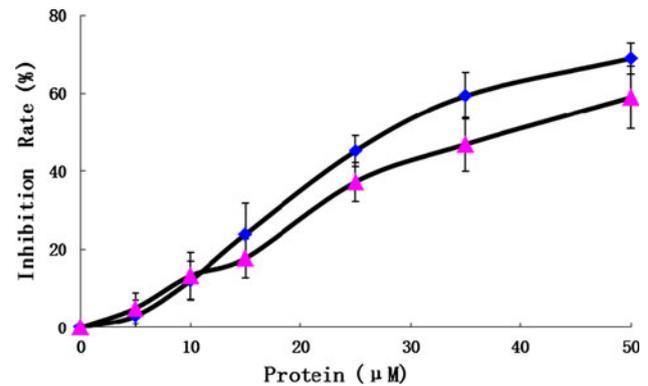
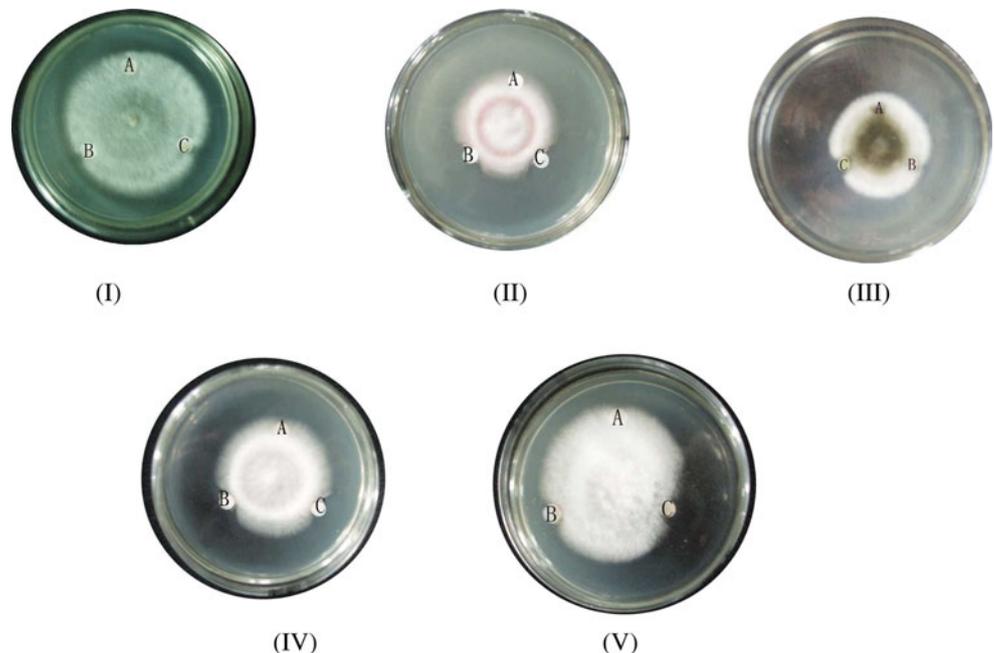


Fig. 9 Inhibitory activity of trypsin inhibitor *Glytine* toward tumor cells BEL-7402 and SHSY5Y (circle and triangle represent SHSY5Y and BEL-7402, respectively). Each data point represents the mean value of triplicate determinations

surprisingly, whey protein showed no significant antiproliferative activity on the cells mentioned above (data not shown).

Significantly, *Glytine* possessed antiproliferative activities against cancer cells in addition to its antifungal activities. Some other antifungal proteins have also been reported to demonstrate antibacterial activities [15, 23–25] as well as antiproliferative activities against tumor cell lines [20, 23, 24, 26, 27]. Our investigation represents a significant addition when compared with those previous studies.

Conclusion

Glytine, a trypsin–chymotrypsin inhibitor with antiproliferative activity against cancer cells and antifungal potency

against several fungal species, has been investigated for the first time in this study. The seeds of the Chinese black soybean *Glycine max* (L.) Merr contain a trypsin–chymotrypsin inhibitor with versatile biological activities, suggesting an exploitable potential for this class of antifungal protein in the food industry and agriculture, as well as in the field of medical treatment.

Acknowledgments This work was supported by the National Natural Science Foundation of China (No. 31071498), Project for Excellent Talents in New Century of Fujian Province (No. JA10012), The S&T projects of Fujian Provincial Science & Technology Hall (No. 2012N0015 & 2012S0053), The S&T development fund of Fuzhou University (No. 2010-XQ-18), The S&T projects of Fujian Provincial Department of Education (No. JA11019), and National Basic Research Program of China (No. 2010CB530605). The authors are very grateful to Dr. Alastair Macdonald from the University of Edinburgh, UK, for his correction of this draft.

Conflict of interest None.

Compliance with Ethics Requirements This article does not contain any studies with human or animal subjects.

References

- Ng TB, Huang B, Fong WP, Yeung HW (1997) *Life Sci* 61:933–949
- Wang HX, Ng TB (2006) *Biochem Biophys Res Commun* 342:349–353
- Birk Y (2003) In: *Plant protease inhibitors*. Berlin: Springer
- Kennedy AR (1995) *J Nutr* 125:733–743
- Kobayashi H, Suzuki M, Tanaka Y, Kanayama N, Terao T (2003) *J Biol Chem* 278:7790–7799
- Cavalcanti MDM, Oliva MLV, Fritz H, Jochum M, Mentele R, Sampaio M, Coelho LCBB, Batista IFC, Sampaio CAM (2002) *Biochem Biophys Res Commun* 291:635–639
- Kumar P, Rao AG, Hariharaputran S, Chandra N, Gowda LR (2004) *J Biol Chem* 279:30425–30432
- Zhao M, Naude RJ, Muramoto K, Oelofsen W (1996) *Int J Pept Protein Res* 48:174–181
- Deshimaru M, Hanamoto R, Kusano C, Yoshimi S, Terada S (2002) *Biosci Biotechnol Biochem* 66:1897–1903
- Wong RC, Fong WP, Ng TB (2004) *Peptides* 25:163–169
- Murdock LL, Huesing JE, Nielsen SS, Prat RC, Shade RE (1990) *Phytochem* 29:85–89
- Wang SY, Ng TB, Chen T, Lin DY, Rao PF, Ye XY (2005) *Biochem Biophys Res Commun* 327:820–827
- Wang SY, Lin J, Ye MY, Ng TB, Rao PF, Ye XY (2006) *Peptides* 27:3129–3136
- Ng TB (2004) *Peptides* 25:1215–1222
- Wang SY, Wu JH, Ng TB, Ye XY, Rao PF (2004) *Peptides* 25:1235–1242
- Lowry OH, Rosebrough NJ, Farr AL, Randall RJ (1951) *J Biol Chem* 193:265–275
- Laemmli UK, Favre M (1973) *J Mol Biol* 80:575–599
- Tanaka S, Takikawa H, Wakabayashi K (1981) *Endocrinol Jpn* 28:335–345
- Wang SY, Rao PF (2010) *Eur Food Res Technol* 231:331–338
- Lin P, Ng TB (2008) *Process Biochem* 43:992–998
- Raj PA, Dentino AR (2002) *FEMS Microbiol Lett* 266:9–18
- Shao B, Wang SY, Zhou JW, Ke LL, Rao PF (2011) *Process Biochem* 46:1554–1559
- Wang SY, Shao B, Rao PF, Lee YY, Ye XY (2007) *J Agric Food Chem* 55:9792–9799
- Mastrolorenzo A, Rusconi S, Scozzafava A, Barbaro G, Supuran CT (2007) *Curr Med Chem* 14:2734–2748
- Flores T, Alape-Giron A, Flores-Diaz M, Flores HE (2002) *Plant Physiol* 128:1291–1302
- Qureshi A, Colin PL, Faulkner DJ (2000) *Tetrahedron* 56:3679–3685
- Wang SY, Rao PF, Ye XY (2009) *Appl Microbiol Biotech* 82:79–86