

# Soluble expression, purification, and characterization of *Gloydius shedaoensis* venom glosedobin in *Escherichia coli* by using fusion partners

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**Abstract** Glosedobin, a thrombin-like enzyme from the venom of *Gloydius shedaoensis*, is usually produced as inclusion bodies in *Escherichia coli* cell. In this work, glosedobin was separately fused with three fusion partners NusA, GST, and TrxA at its N terminus and then was expressed as fusion proteins in *E. coli*. The results showed that the NusA was the most efficient fusion partner to improve the solubility of recombinant glosedobin. The purified NusA-fused glosedobin with an overall yield of 64.6% was resolved as one band in the SDS-PAGE gel with molecular mass of about 90 kDa. Both fibrinogen clotting and fibrinogenolytic activities were found for the recombinant product. The purified NusA-fused glosedobin exhibited amidolytic activity of 506 U/mg under optimal conditions of pH of 8.0 and 40°C. The inhibition study of NusA-fused glosedobin by various inhibitors showed that serine protease inhibitors, phenylmethylsulphonyl fluoride, and *N*-tosyl-L-phenylalanine chloromethyl ketone, strongly inhibited its amidolytic activity, whereas ethylenediaminetetraacetic acid as well as heparin and hirudin did not, suggesting that NusA-fused glosedobin exhibited the same characteristics as the native form of glosedobin. The strategy of this work may contribute to improve the soluble expression level of other thrombin-like enzymes from snake venom in *E. coli*.

**Keywords** Fibrinolytic enzyme · GST · NusA · TrxA · Snake venom · Thrombin-like enzyme

## Introduction

Snake venoms are very complex mixtures containing a wide variety of biologically active proteins and polypeptides (Lu et al. 2005; Markland 1998). Thrombin-like enzymes from snake venom belong to the serine protease family, exhibiting fibrinogenolytic activity that cleave the A $\alpha$ -chain or the B $\beta$ -chain of fibrinogen to release fibrinopeptide A or fibrinopeptide B, respectively (Castro et al. 2004). Since thrombin-like enzymes do not activate factor XIII, the resulting fibrin clot is not cross-linked and easily removed from circulation by the reticuloendothelial system or normal fibrinolytic system (Kini et al. 2001). Because of these physiological properties, thrombin-like enzymes have been used as antithrombotic agents for the treatment of occlusive thrombi (Koh et al. 2006; Marsh and Williams 2005).

To date, the amino acid sequences of dozens of thrombin-like enzymes have been determined partially or completely from various snake species (Castro et al. 2004; Pirkle 1998). Glosedobin (formerly named defibrase) is a thrombin-like enzyme obtained from the venom of *Gloydius shedaoensis* and has been cloned and partially characterized in our previous work (Yang et al. 2002). Glosedobin has high physiological activities of thrombin-like enzymes and, therefore, has been applied in the treatment of the thrombotic diseases. However, similar to other thrombin-like enzymes such as acutin from *Agkistrodon acutus* and pallabin from *Agkistrodon halys pallas*, glosedobin was also expressed as biologically inactive inclusion bodies in *E. coli* (Pan et al. 1999; Fan et al. 1999). This may be attributed to that it is a cytotoxic protein and contains 12 cysteine residues forming six disulfide bonds (Yang et al. 2003). Many efforts have been attempted to improve the soluble expression level of recombinant thrombin-like enzymes in various microorganisms. Habutobin was expressed in a baculoviral system (Sunagawa et al. 2007). Soluble form of

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gloshedobin has been successfully produced in the yeast *Pichia pastoris* (Yang et al. 2002). However, the solubility of synthesized protein has not yet reached a satisfactory level.

Although a number of expression systems are available for production of proteins of therapeutic or commercial interest, *E. coli* has always been the first choice as expression host because of its low cost, fast growth, and easy operation (Baneyx 1999; Jana and Deb 2005). Many efforts such as reducing cultivation temperature (Vasina and Baneyx 1997), co-expressing of molecular chaperones (de Marco and De Marco 2004), and altering *E. coli* expression strain (Miroux and Walker 1996) have been attempted to overcome the difficulty in generating soluble recombinant protein in *E. coli*. Fusion protein expression by cloning a unique fusion tag into the N-terminal of a target protein is a preferable strategy for enhancing the solubility of recombinant proteins in *E. coli* (Esposito and Chatterjee 2006). Commercial fusion tags such as thioredoxin (TrxA, LaVallie et al. 1993), glutathione-S-transferase (GST, Nygren et al. 1994), N-utilization substance A (NusA, Nallamsetty and Waugh 2006), and disulfide oxidoreductase (DsbA, Zhang et al. 1998b) have been revealed to either increase the expression level or improve solubility of recombinant protein.

In this study, three soluble partners (NusA, GST, and TrxA) fused at N terminus of gloshedobin were expressed in *E. coli*, and the efficiency to increase the solubility of gloshedobin was compared. It was found that NusA was the most efficient fusion partner. The purified NusA-fused gloshedobin exhibited satisfactory enzymatic properties including amidolytic, fibrinogen clotting, and fibrinolytic activities.

## Materials and methods

### Host strains, plasmids, and enzymes

*E. coli* JM109 (Takara, Dalian, China) was used as the host for DNA manipulation. *E. coli* BL21-Gold (DE3; Stratagene,

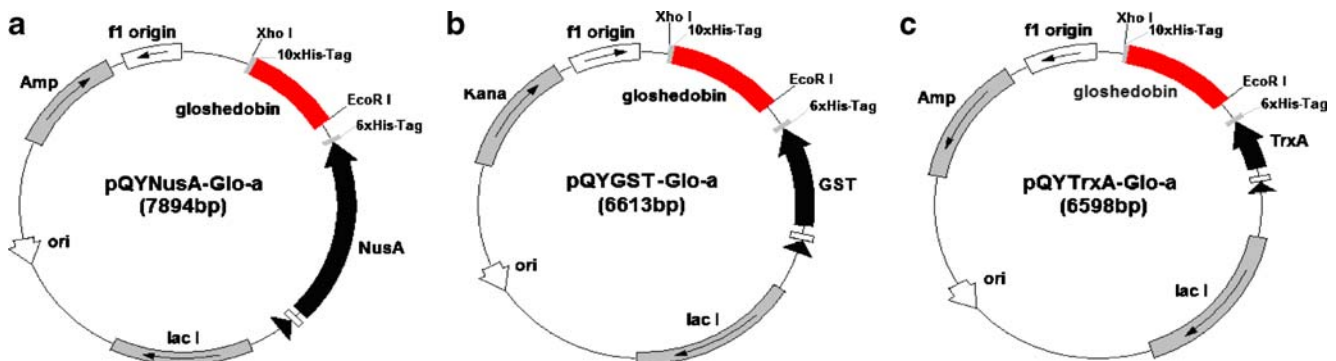
USA) was used as the host for expression of recombinant protein. pMD18-T Simple Vector (Takara, Dalian, China) was used as cloning vector. Plasmids pET-43.1a, pET-42a, and pET-32a (Novagen, Madison, WI, USA) were used to construct expression vectors. T4 DNA ligase, *Ex Taq* DNA polymerase, and all the restriction enzymes were obtained from Takara (Dalian, China).

### Construction of cloning and expression vectors

The gene-encoding gloshedobin was amplified from plasmid pPIC9K-TLE (Yang et al. 2002) by polymerase chain reaction (PCR) with the N-terminal primer 5'-CGGAATTC ATCATTGGAGGTGATGAA -3' (*EcoRI* site is underlined) and the C-terminal primer 5'-CCGCTCGAGTTAGTG GTGGTGGTGGTGG TGGTGGTGGTGGTGGGG GGCAGGTTGCATC-3' (*XhoI* site is underlined). The PCR product was digested with *XhoI-EcoRI* and cloned into the *XhoI-EcoRI* site of the pMD18-T simple vector and then transformed into *E. coli* JM 109. Nucleic acid sequences of the cloning DNA fragment were confirmed by DNA sequencing (BigDye™ Kit, Applied Biosystems, USA) and ABI PRISM™ 3730XL DNA Analyzer, according to the recommended protocols. The target DNA fragment was further subcloned in the same site of pET-43.1a, pET-42a, and pET-32a vectors, resulting in pQYNusA-Glo-a, pQYGST-Glo-a, and pQYTrxA-Glo-a, respectively (Fig. 1). The resulting vectors were transformed into *E. coli* BL21-Gold (DE3) for fusion protein expression.

### Expression of gloshedobin fusion proteins

The single colony of *E. coli* BL21-Gold (DE3) harboring the expression vector was inoculated in 30 ml of Luria–Bertani medium containing 100 µg/ml ampicillin and then cultivated at 37°C until the optical density (OD<sub>600</sub>) reaches 0.6. The cells were harvested by centrifugation at 4,000×g for 10 min and resuspended in 3 l fresh Terrific Broth



**Fig. 1** Schematics of three expression constructs of fusion proteins. The gene encoding gloshedobin was inserted between *XhoI* and *EcoRI*. **a** pQYNusA-Glo-a containing T7 promoter and NusA tag; **b**

pQYGST-Glo-a containing T7 promoter and GST tag; **c** pQYTrxA-Glo-a containing T7 promoter and TrxA tag

medium containing 100 µg/ml ampicillin and 1% (*w/v*) glucose. Expression of the fusion protein was induced with isopropyl-β-D-thiogalactoside (IPTG) at a final concentration of 1.0 mM, and the culture was incubated for an additional 8 h at 25°C. The cells were harvested by centrifugation at 8,000×*g* for 15 min and washed with TE buffer (20 mM Tris-HCl, 10 mM ethylenediaminetetraacetic acid (EDTA), pH8.0). Approximated 30 g (wet weight) cells were obtained from 3 l culture. After centrifugation, the cell pellets was resuspended in 40 ml (for 1 l culture) ice-cold extraction buffer (20 mM Tris-HCl, 100 mM NaCl, 1 mM EDTA, pH 8.0, 0.5 % (*w/v*) glycerol) and lysed by ultrasonication at ice-cold temperature using UP400S instrument (Dr. Hielscher GmbH, Stuttgart, Germany). The cell lysis was centrifuged at 12,000 rpm for 15 min to separate soluble (supernatant) and precipitated (pellet) fractions.

#### Purification of glosedobin fusion protein

Solid ammonium sulfate was added into the supernatant to reach 1 M. After centrifugation at 12,000 rpm for 20 min at 4°C, the supernatant was loaded onto a Phenyl Sepharose FF column (GE Healthcare-Amersham Biosciences) followed by elution with a gradient of 1–0 M ammonium sulfate. Active fractions were collected and loaded onto a Q-Sepharose HP column (GE Healthcare-Amersham Biosciences) followed by a Mono Q column (GE Healthcare-Amersham Biosciences), eluting by a gradient of 0.2–0.4 and 0.15–0.4 M of NaCl, respectively. The active fractions were concentrated by ultracentrifuge (Amicon Ultra-15, Millipore, USA) and loaded onto a Superdex 200 column (GE Health-Amersham Biosciences) by eluting with elution buffer (50 mM Tris-HCl, 150 mM NaCl, pH7.5). Aliquots of the resulting purified protein were frozen at –20°C for subsequent analysis. Enzymatic activity assay by using substrate *N*α-benzoyl-DL-arginine-*p*-nitroanilide (BAPNA) was applied to monitor the elution of active fractions during every purification step.

#### SDS-PAGE and western blot analysis

SDS-PAGE analysis was performed using 10% resolving gel and 5% stacking gels. The protein bands were visualized by Coomassie brilliant blue R-250 and then analyzed by image-density analysis software (Gel-Pro, USA). Protein samples were subjected to SDS-PAGE and transferred to polyvinylidene difluoride membrane (Millipore, USA). The membranes were blocked with 5% defatted milk at room temperature for 60 min and then incubated with anti-snake venom horse serum (1:500, Shanghai Institute of Biologic Products, Shanghai, China) at room temperature for 3 h. The membranes were washed five times with TBS-T buffer (20 mM Tris-HCl, pH8.0, 150 mM NaCl, 0.15% (*v/v*) Tween-20) and then incubated with horseradish peroxidase-

conjugated rabbit anti-horse IgG (1:5,000, Shanghai Institute of Biologic Products, Shanghai, China) at room temperature for 60 min. Finally, the membranes were washed five times as above, and a 3'-diaminobenzidine kit was used for color development detected (Sigma, St. Louis, Mo, USA).

#### Amidolytic activity assay

Amidolytic activity was determined by incubating 195 µl of the substrate, 2 mM BAPNA (20 mM Tris-HCl, pH7.5) with 5 µl of glosedobin (1.0 µg) in a 96-well plate. The amount of *p*-nitroaniline released was determined by measuring the change in absorbance at 405 nm using a Sinrise Spectrophotometer (Tecan, Austria). One unit of amidolytic activity was defined as the quantity of enzyme required for hydrolyzing 1.0 µM/min BAPNA at 25°C.

#### Fibrinogen clotting and fibrinolytic activity assay

Fibrinogen clotting activity was determined by incubating 100 µl of purified NusA-fused glosedobin with 200 µl of fibrinogen (4 mg/ml) in 20 mM Tris-HCl (pH7.5) containing 150 mM NaCl at 37°C for 6 h in glass tube, and then the occurrence of the fibrinogen clotting phenomenon was verified by photos. The fibrinogen from bovine, purchased from Sigma, is plasminogen-free and of electrophoretical-grade purity.

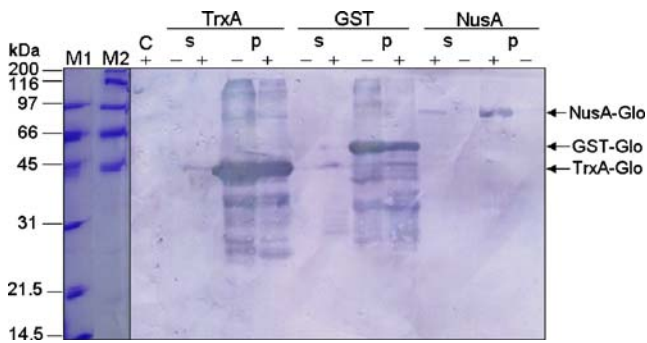
Fibrinolytic activity was determined by incubating 25 µl of purified NusA-fused glosedobin (1.0 µg) with 100 µl of fibrinogen (0.2 mg/ml) in 20 mM Tris-HCl (pH7.8) containing 150 mM NaCl at 37°C for 0.5, 1, 1.5, 2, 2.5, 4, 6, 8, 10, 12, 19, and 24 h, respectively. Then, the reaction mixture was separated by 10% SDS-PAGE to examine the cleavage pattern of fibrinogen chains digested by glosedobin.

#### Effects of pH and temperature on amidolytic activity

The effect of pH on amidolytic activity was determined by preincubating 1.0 µg of enzyme in citrate buffer (20 mM, pH4–6) or Tris-HCl buffer (20 mM, pH7–10) at 25°C for 20 min. The reaction was initiated by addition of 2 mM substrate solution (BAPNA, 20 mM Tris-HCl, pH7.5). The effect of temperature on amidolytic activity was determined by preincubating 1.0 µg of enzyme in 20 mM Tris-HCl (pH 7.5) at different temperatures (20°C, 30°C, 40°C, 50°C, and 60°C) for 20 min. The reaction was initiated by addition of 2 mM substrate solution (BAPNA, 20 mM Tris-HCl, pH7.5).

#### Effect of enzyme inhibitors on the amidolytic activity

The effect of enzyme inhibitors on the amidolytic activity was determined by preincubating 1.0 µg of enzyme in 20 mM Tris-HCl (pH7.5) with inhibitors including 0.1 or



**Fig. 2** Western blot analysis of the recombinant glosedobin fused with soluble fusion tags. *Lane S (+)* represents soluble fraction of cell lysates after IPTG induction; *Lane S (-)* represents soluble fraction of cell lysates before IPTG induction; *Lane P (+)* represents precipitates after IPTG induction; *Lane P (-)* represents precipitates before IPTG induction; *Lane C* host strain alone; *Lane M1, M2* protein molecular weight marker

1 mM phenylmethylsulphonyl fluoride (PMSF), 0.1 or 1 mM *N*-tosyl-L-phenylalanine chloromethyl ketone (TPCK), 1 mg/ml heparin, 0.4 mg/ml hirudin, 3 mM dithiothreitol (DTT), 3 mM  $\beta$ -mercaptoethanol ( $\beta$ -ME), 10 mM EDTA at 25°C for 20 min. The reaction was initiated by addition of 2 mM substrate solution (BAPNA, 20 mM Tris-HCl, pH7.5).

## Results

### Construction of the expression vectors

Glosedobin gene was successfully amplified from the plasmid pPIC9K-TLE and ligated into plasmid pET-43.1a, pET-42a, and pET-32a vectors, resulting in pQYNusA-Glo-a, pQYGST-Glo-a, and pQYTrxA-Glo-a, respectively. In these recombinant plasmids, glosedobin was fused with

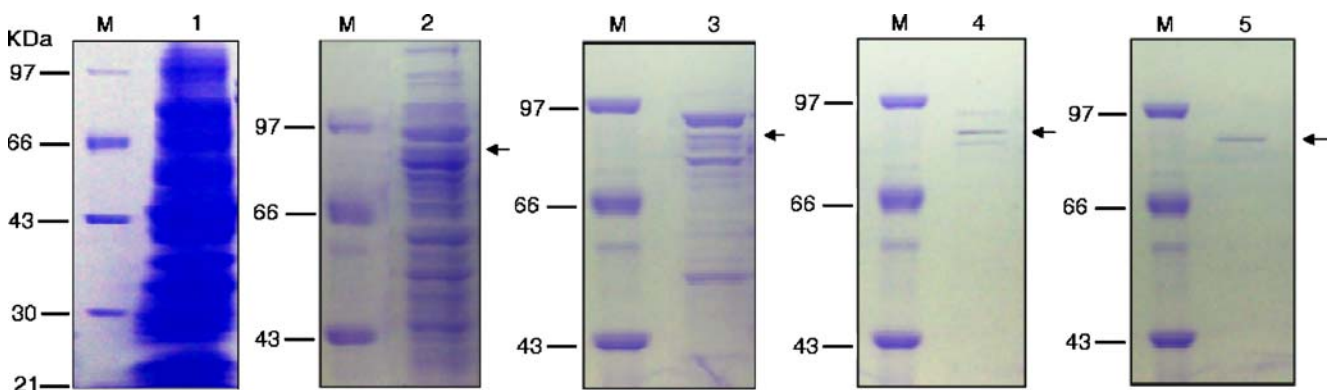
fusion partners NusA, GST, and TrxA, respectively, under the control of T7 promoter (Fig. 1). All constructs were confirmed by DNA sequencing.

### Expression of NusA-, GST-, and TrxA-fused glosedobin

The fusion expression vectors (pQYTrxA-Glo-a, pQYGST-Glo-a, and pQYNusA-Glo-a) encoding the target protein combined with the soluble tags were transformed into *E. coli* BL21-Gold (DE3) to express proteins of interest, respectively. When the culture was propagated at 37°C until the optical density ( $OD_{600}$ ) reaches 0.6, the expression of the fusion protein was induced with 1.0 mM of IPTG, and then the culture was grown for an additional 8 h at 25°C. Consequently, approximated 10 g (wet weight) cells were obtained from 1 l culture. After lysed by ultrasonication, the supernatant and pellet fractions of fusion proteins were analyzed by western blotting using horse anti-snake venom antibody. As shown in Fig. 2, although TrxA-, GST-, and NusA-fused glosedobins appeared in soluble fraction of the cell lysate after induced with IPTG-, TrxA-, and GST-fused glosedobin were mainly in the form of inclusion bodies in *E. coli*. Moreover, TrxA- and GST-fused glosedobin were also expressed even without IPTG induction, which was also seen in Fig. 2. Unlike these two, NusA-fused glosedobin was only produced after the induction of IPTG and mainly in the soluble form in the cytosol of *E. coli*. Therefore, NusA-fused glosedobin was selected as example in the subsequent experiments including.

### Purification of NusA-fused glosedobin

NusA-fused glosedobin was purified in this study by four chromatographic steps: hydrophobic interaction chromatography, two anion exchanges chromatographies, and gel



**Fig. 3** SDS-PAGE analysis of purified NusA-fused glosedobin by four-step chromatographies. *Lane M* protein molecular weight marker; *lane 1* cell lysates; *lane 2* the eluent from hydrophobic interaction chromatography using Phenyl Sepharose FF column; *lane 3* the eluent

from anion exchanges chromatography using Q-Sepharose HP column; *lane 4* the eluent from anion exchange chromatography using Mono Q column; *lane 5* the eluent from gel filtration chromatography using Superdex 200 column



filtration chromatography. Specific activity of NusA-fused glosheedobin was determined by its amidolytic activity by using the substrate, BApNA. As shown in Fig. 3, SDS-PAGE showed that the purified NusA-fused glosheedobin was observed as a single band in the gel with molecular masses of about 90 kDa. After these four-step chromatographies, the specific activity and overall recovery of purified NusA-fused glosheedobin was 506 U/mg and 64.6%, respectively (Table 1). The recovery and purification factors of each chromatography step are summarized in Table 1.

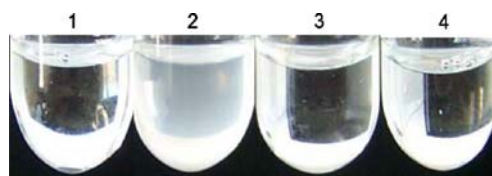
#### Fibrinogen clotting and fibrinolytic activities of purified NusA-fused glosheedobin

In vitro, thrombin-like enzymes are serine protease, which resemble thrombin, play a key role in coagulation because they cleave fibrinogen to cross-linked fibrin by releasing either fibrinopeptide A or fibrinopeptide B. Therefore, thrombin-like enzymes exhibit fibrinogen clotting and fibrinolytic activities in vitro. However, in vivo thrombin-like enzyme different from thrombin do not activate factor XIII. Consequently, fibrin cleaved by these proteases cannot be cross-linked, and then easily removed from circulation by the reticuloendothelial system or normal fibrinolytic system.

In the case of fibrinogen clotting activity assay, the fibrinogen was incubated with purified NusA-fused glosheedobin at 37°C for 6 h in glass tube, and then the occurrence of the fibrinogen clotting phenomenon was verified by photos. As shown in Fig. 4, the solution containing fibrinogen and purified NusA-fused glosheedobin was turbid; however, other glass tube was clear. These results indicated that the addition of purified NusA-fused glosheedobin only led to the formation of a clotting phenomenon. In the case of fibrinolytic activity assay, the degradation products of fibrinogen digested with NusA-fused glosheedobin were analyzed by SDS-PAGE. As shown in Fig. 5, the band of  $\alpha$  chain of fibrinogen became invisible on the gel after 2 h of the reaction. Moreover, the  $\beta$  chain of fibrinogen was fully digested after 24 h of the reaction. However, the remaining  $\gamma$  chain of fibrinogen seems to be stable, since the band of  $\gamma$  chain was intact even after 24 h. These results indicated that NusA-fused glosheedobin exhibited the thrombin-like activities.

**Table 1** Recoveries and purification of NusA-fused glosheedobin

Purification step	Specific activity (U/mg)	Purification factor	Yield (%)
Lysate supernatant	0.163	1	100
Phenyl Sepharose FF column	8.20	50	84.9
Q Sepharose HP column	95.32	585	75.5
Mono Q column	343.13	2,105	70.1
Superdex 200 column	506.00	3,104	64.6



**Fig. 4** Fibrinogen clotting activity of purified NusA-fused glosheedobin; 1 0.4 mg/ml of fibrinogen only; 2 0.4 mg/ml of fibrinogen with purified NusA-fused glosheedobin; 3 0.4 mg/ml of fibrinogen with concentrated culture medium after IPTG induction; 4 0.4 mg/ml of fibrinogen with cytosol of *E. coli* carrying empty plasmid after IPTG induction

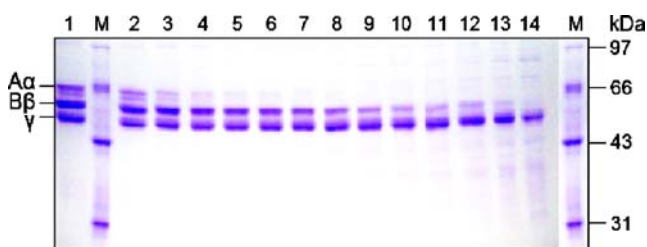
#### Effects of pH, temperature, and inhibitors on amidolytic activity

The amidolytic activity of the purified NusA-fused glosheedobin was determined using BApNA as substrate. As shown in Fig. 6, the optimum conditions for the enzyme's amidolytic activity were determined to be 40°C and pH8.0 (Fig. 6a, b).

The effect of various inhibitors on the amidolytic activity of NusA-fused glosheedobin was evaluated. As shown in Table 2, the concentration of 1 mM of PMSF completely inhibited this activity. The amidolytic activity of NusA-fused glosheedobin was partially affected by TPCK, DTT, and  $\beta$ -ME. However, heparin and EDTA had no effect on this activity.

#### Discussion

The expression of recombinant thrombin-like enzymes in microorganisms is still of research interest because the amount of thrombin-like enzymes extracted from snake venom is very limited. Some thrombin-like enzymes have been successfully synthesized in various microorganisms by recombinant DNA technology. Salmobin, a thrombin-like enzyme obtained from *Agkistrodon halys*, was expressed by cell surface display on *E. coli* using ice nucleation protein (Jeong et al. 2001). In addition, soluble form of batroxobin and ancrod has successfully been obtained in the yeast strain *P. pastoris* (You et al. 2004; Yu et al. 2007). However, the expression level of proteins synthesized by these systems could not



**Fig. 5** SDS-PAGE analysis of fibrinogenolytic activity of purified NusA-fused glosheedobin. Lane 1 undigested fibrinogen; lanes 2–14 fibrinogen solution was incubated with purified NusA-fused glosheedobin at 37°C for 0, 0.5, 1, 1.5, 2, 2.5, 4, 6, 8, 10, 12, 19, and 24 h, respectively

satisfy requirements of biochemical analysis, therapeutics, or structural studies.

*E. coli* is the preferred host cell for the expression of recombinant proteins because of its well-characterized genetics, rapid growth, and the availability of numerous vectors (Baneyx 1999; Jana and Deb 2005). However, thrombin-like enzymes are difficult to be expressed in the soluble form in *E. coli* because they are cytotoxic and single-chain cystein-rich proteins. Correct folding of recombinant thrombin-like enzymes in term of enzymatic activities is still a big challenge.

Calobin, a thrombin-like enzyme obtained from *Agkistrodon caliginous*, was co-expressed with disulfide isomerase (DsbC) and TrxA (Yuan et al. 2004). The free form of TrxA and DsbC were efficient for the soluble expression of calobin. However, the recombinant calobin-TrxA-DsbC only retained the fibrinogenolytic activity but lost most of amidolytic activity and even all the fibrinogen clotting activity. Therefore, in this work, *NusA*, *GST*, and *TrxA* genes were expressed as fusion partners, not in the free form. As shown in Fig. 2, although NusA, GST, and TrxA were all efficient for the soluble expression of glosheedobin, the solubility of NusA-fused glosheedobin was the most satisfactory one. In addition, the expression vectors, pQYTrxA-Glo-a and pQYGST-Glo-a, failed to be strictly controlled under T7 promoter because recombinant proteins were observed to be produced even without IPTG induction (Fig. 2). These results indicated that NusA was the most

**Table 2** Effects of various inhibitors on the amidolytic activity of NusA-fused glosheedobin

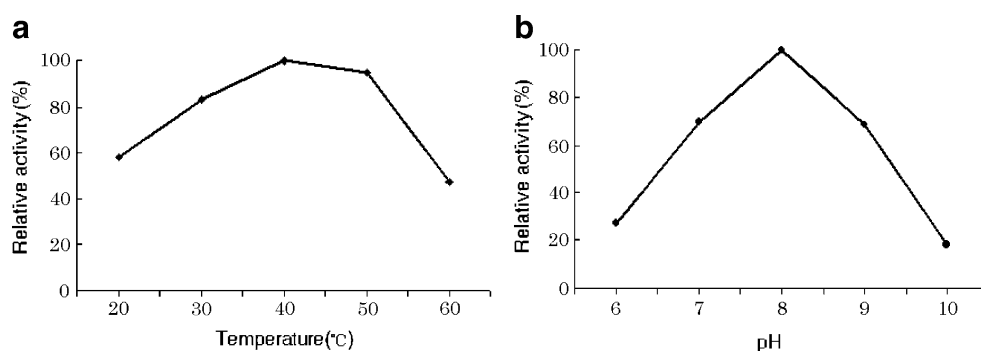
Inhibitors	Concentration	Relative activity (%)
Control		100
PMSF	0.1 mM	50
PMSF	1 mM	0
TPCK	0.1 mM	61
TPCK	1 mM	23
Heparin	1 mg/ml	97
Hirudin	0.4 mg/ml	87
DTT	3 mM	85
$\beta$ -ME	3 mM	86
EDTA	10 mM	97

efficient fusion partner for improving the soluble expression level of glosheedobin.

During molecular cloning, his-tag at C terminus of recombinant glosheedobin was designed in an attempt to simplify the purification procedure. Although his-tag purification strategy is widely applied in the simplification of purification of recombinant proteins, it is unfortunate that most glosheedobin existed in unbound fractions, which was observed by enzymatic activity assay. Similarly, it is noted for its ineffective binding to Ni-chelating ligands (Yuan et al. 2004; Debeljak et al. 2006). The reason may not be clear until detailed structural information is revealed. Therefore, in this work, four chromatographic steps were used to purify the NusA-fused glosheedobin, which contained hydrophobic interaction chromatography, two anion exchanges chromatographies, and gel filtration chromatography. According to SDS-PAGE analysis shown in Fig. 3, the nonspecific bands gradually disappeared, and the NusA-fused glosheedobin finally became a single band. These results indicated that although four chromatography steps were cumbersome, they were very effective.

The fibrinogenolytic activity is a main physiological property of thrombin-like enzymes in terms of their potential clinical application in thrombosis and anticoagulation. In this work, the purified NusA-fused glosheedobin

**Fig. 6** Effects of different pH and temperatures on amidolytic activity of purified NusA-fused glosheedobin. Enzyme activity was performed at different temperatures (a) and pH (b) using *N* $\alpha$ -benzoyl-DL-arginine-*p*-nitroanilide as substrate



preferentially digested the A $\alpha$  chain then slowly cleaved the B $\beta$  chain, however, did not affect the  $\gamma$  chain (Fig. 5). This is very similar to most of thrombin-like enzymes (Nishida et al. 1994; Zhang et al. 1998a), although other individual thrombin-like enzymes which preferentially degraded the A $\alpha$  or B $\beta$  chains (Sant'Ana et al. 2008) or degraded both A $\alpha$  and B $\beta$  chains with similar efficiency, were reported (Castro et al. 2004). In addition, the purified NusA-fused glosedobin in this work also exhibited fibrinogen clotting activity and amidolytic activity, suggesting that NusA fusion tag did not affect the thrombin-like enzyme characteristics of glosedobin.

The optimum conditions for the enzyme's amidolytic activity were determined to be 40°C and pH8.0, similar to that of other thrombin-like enzymes. That serine protease inhibitors PMSF (1 mM) and TPCK (1 mM) inhibited the amidolytic activity of NusA-fused glosedobin by 100% and 77%, respectively (Table 2), is consistent with the fact that it belongs to the serine protease family. The chelator EDTA did not affect its amidolytic activity, also confirming that the recombinant glosedobin was a serine protease.

High concentration of DTT (10 mM) and  $\beta$ -ME (10 mM) had little effect on the amidolytic activity of the recombinant glosedobin in this study, suggesting that recombinant glosedobin was a relatively stable protein in comparison with other thrombin-like enzymes. Bothrombin (Nishida et al. 1994) and stejnobin (Zhang et al. 1998a) were reported to be significantly inhibited by DTT and  $\beta$ -ME. However, like glosedobin, harobin, a fibrinogenolytic serine protease from the sea snake *Lapemis hardwickii* (He et al. 2007), was reported to be resistant to 10 mM DTT. Heparin, a natural thrombin inhibitor, did not affect the amidolytic activity, suggesting that the recombinant glosedobin differed from thrombin in structure. From a clinical point of view, it is important that heparin as well as hirudin does not inhibit its enzymatic activity, which indicates that glosedobin could combine with heparin or hirudin in the treatment of thromboembolic and cardiopulmonary diseases.

In conclusion, we investigated the soluble expression of glosedobin fused with three soluble tags and found that the solubility of NusA-fused glosedobin was the best. The purified NusA-fused glosedobin exhibited satisfactory thrombin-like enzyme activities including amidolytic activity, fibrinogen clotting activity, and fibrinogenolytic activity. This study provided a method useful for the preparation of other recombinant thrombin-like enzymes of clinical interest.

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