



Recombinant expression and purification of the bovine acidic Seminal Fluid Protein

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Abstract

The acidic Seminal Fluid Protein (aSFP), a 12.9 kDa protein is a marker for bovine semen freezability possibly due to its antioxidant activity and effect on sperm mitochondrial function. However, its precise function on sperm preservation during freezing-thaw is poorly understood. The use of recombinant DNA technology allows new approaches on the study of function and structure of proteins, and its production in prokaryote systems offers several advantages. The present work describes the recombinant expression of the bovine aSFP and its binding properties. A cDNA library from the bovine seminal vesicle was used as template for amplification of the aSFP coding region. The amplicon was cloned into a pET23a(+) vector and transformed into *E.coli* BL21 pLysS strain. The recombinant expression was obtained in *E. coli*. One step ion immobilized affinity chromatography was performed, resulting in high yield of purified protein. To determine the bioactivity of the raSFP, the protein was incubated in different concentrations with 10⁷ spermatozoa at 37°C for 5 h. Western blotting and fluorescence microscopy analyses showed the ability of the recombinant aSFP to attach to the spermatozoa. Based on our results, the described method can be used to obtain mg levels of recombinant aSFP.

Keywords: bovine, recombinant protein, seminal plasma, spermatozoa.

Introduction

Seminal plasma (SP), a physiological secretion from multiple glands of the male reproductive tract, is a very complex fluid and is the natural medium for final maturation of spermatozoa through hormonal, enzymatic and surface-modifying events (Mann and Lutwak-Mann, 1981). The proteomic approach of seminal plasma is a valuable tool to identify proteins playing key roles on sperm maturation, viability and fertilization. Although the identification of SP proteins (Assumpção *et al.*, 2005; Kelly *et al.*, 2006; Moura *et al.*, 2006, 2007, 2010) provided many insights about their correlation with biological phenomena, most of

their precise functions are still unknown. Fertility-associated proteins were described in this fluid, as well as glycoproteins which participate in sperm-egg recognition, modulation of spermatozoa capacitation and the acrosome reaction (Killian *et al.*, 1993; Therien *et al.*, 1995; Calvete *et al.*, 1996; Iborra *et al.*, 1996).

The contribution of SP proteins to freezability is also a matter of study in several species, including bull (Roncoletta *et al.*, 2000; Jobim *et al.*, 2004; Bergeron *et al.*, 2007; Manjunath *et al.*, 2007) ram (Barrios *et al.*, 2000; Rebolledo *et al.*, 2007), buffalo (Harshan *et al.*, 2006; Asadpour *et al.*, 2007) and boar (Casas *et al.*, 2009). Comparing the SP protein profile from bulls with high and low semen freezability, Jobim *et al.*, (2004) suggested the acidic Seminal Fluid Protein (aSFP) as a marker of good semen freezability. The 12.9 kDa aSFP is an acidic nonglycosylated protein (pI 4.8; Einspanier *et al.*, 1991), mainly synthesized by the ampulla and seminal vesicle epithelium of the bull in high concentrations (2-7 mg/ml in SP; Einspanier *et al.*, 1993; Dostàlovà *et al.*, 1994) and has recently been described in epididymal fluid (Moura *et al.*, 2010). This protein has not been detected in any other bovine tissues nor in other mammalian species, such as goat, sheep, pig, rat, dog, or human (Wempe *et al.*, 1992; Einspanier *et al.*, 1993). It presents approximately 50% amino acid sequence identity with polypeptides of the spermadhesin family (Einspanier *et al.*, 1994), but with different biological characteristics, aSFP binds only loosely to the acrosomal cap of ejaculated bovine sperm and is quantitatively released during *in vitro* capacitation (Dostàlovà *et al.*, 1994). In addition, aSFP possesses neither carbohydrate nor zona pellucida binding abilities, strongly indicating that aSFP may not be involved in gamete interaction (Dostàlovà *et al.*, 1994; Ekhlasi-Hundrieser *et al.*, 2008).

Recently, the production of recombinant forms of seminal plasma proteins has been adopted as a new tool for understanding the biological activities of spermadhesins (Ekhlasi-Hundrieser *et al.*, 2008; Cajazeiras *et al.*, 2009). Considering its possible role on semen freezability, the aim of this work was to purify a bioactive recombinant form of aSFP and to study its binding properties on bovine semen.

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Materials and Methods

Materials

Expression vector pET23a(+), as well as *Escherichia coli* strains BL21 (DE3) pLysS, BL21 (DE3) were from Novagen (EMD Biosciences, La Jolla, CA, USA); JM109 strains were from Promega (Madison, WI, USA). TOPO TA kit, all modification enzymes, antibody anti His-tag, secondary anti mouse and *E. coli* strains BL21 (DE3) Star and BL21 SI were from Invitrogen (Carlsbad, CA, USA). RNA extraction kit, Miniprep DNA isolation kit and Miniprep Plasmid DNA purification kit were from Macherey-Nagel (Düren, Germany). Reagents to detect chemiluminescence (ECL) were purchased from Amersham Pharmacia Biotech (Piscataway, NJ, USA). Hybond-C nitrocellulose membranes were from Hybond™ ECL™ (Hybond™ ECL™ nitrocellulose membrane, Amersham Biosciences, Freiburg, Germany). X-ray films were purchased from Kodak (Kodak X-Omat, Rochester, NY, USA).

Cloning of cDNA sequence into expression vector

For expression of aSFP with N-terminal His-tag, bovine seminal vesicle cDNA was used as a template for polymerase chain reaction (PCR) amplification of aSFP. Synthetic oligonucleotide primers (forward: 5'-**GTCCATATGAAGCTGTCCAGCGTCAT**-3'; reverse: 5'-**AGTCTCGAGAGCTTGTGGATCCT**-3') were designed based on the aSFP sequence available on the GeneBank (accession number NM_174616; Wempe *et al.*, 1992) to amplify the codon region. The 5' *Nde*I and 3' *Xho*I restriction sites are underlined, and the start codon is shown in bold.

PCR amplification was performed in a thermocycler TC-4000 (Techne UK, Staffordshire, UK) using *Taq* DNA polymerase, under the following conditions: 94°C, 2 min; 36 cycles of 94°C, 30 s; 56°C 30 s; 72°C, 1 min, and the final elongation of 72°C for 5 min. The resulted aSFP amplicon (404 bp) was cloned into pCR2.1 vector, excised from this vector using the restriction enzyme pair mentioned above, and ligated into the expression vector pET23(a)+ linearized with the same enzyme pair. Digested insert and vector were run on a 1.5% agarose gel, gel purified using a commercial kit and then incubated overnight at 16°C using T4 DNA ligase. Ligation reactions were transformed into electrocompetent *E. coli* JM109 cells, and plasmid DNA was isolated. The resulting construct pET-aSFP was sequenced using the Big Dye Terminator® v 3.1 Cycle Sequencing Kit (Applied Biosystems, Foster City, CA, USA) and the automated sequencer ABI PRISM® 3100 Genetic analyzer (Applied Biosystems). Dyeing reactions were performed using a *GeneAmp PCR System 9700* (Applied Biosystems) and data analyzed using *Data*

Collection v1.0.1 (Applied Biosystems). The following primers were used for sequence confirmation: forward 5'-TAATACGACTCACTATAGGG-3'; reverse 5'-GCTAGTTATTGCTCAGCGG-3'; forward 5'-CGCCACCTATTACGGACCGAAAAC-3'; reverse 5'-TCCCTAAGACAGGAGATCCTGGCA-3'.

Protein expression in *E. coli*

For protein expression assays, plasmids were transformed into *E. coli* BL21 pLysS competent cells using methods described elsewhere (Sambrook and Russell, 2001). Transformed bacteria were grown overnight on LB-agar plates containing 100 µg/ml ampicillin and/or 50 µg/ml chloramphenicol, after which single colonies were used to inoculate 2x YT medium containing the same antibiotic. For protein expression, 250 ml 2x YT medium was inoculated with 1/100 volume of overnight culture; bacteria were grown at 37°C with shaking at 200 rpm until the OD₆₀₀ reached 0.6 - 0.8 and then the lactose analogue, isopropyl-β-D-thiogalactopyranoside (IPTG) was added to a final concentration of 0.5 mM to induce expression. Three hours after induction, cell cultures were harvested by centrifugation at 5,000 x g for 10 min at 4°C.

Protein purification using immobilized metal affinity chromatography

Cell pellet from 250 ml induced culture was suspended in 10 ml of denaturing lysis buffer (50 mM NaH₂PO₄, 8 M urea, pH 8.0), incubated for 30 min in the presence of lysozyme (1 mg/ml), disrupted by sonication (6 cycles of 10 s on ice), and centrifuged at 20,000 x g for 30 min at 4°C, as described in the Invitrogen protocol handbook. Supernatant was loaded onto an Econopack column (Bio-Rad) filled with 1 mL of Ni-NTA agarose resin (Invitrogen), prepared and equilibrated according to manufacturer's instructions. Washing steps were performed using 5 to 8 bed volumes of denaturing washing buffer (50 mM NaH₂PO₄, 8 M urea) with decreasing pH (pH 6.0 and pH 5.3). Recombinant His-tagged aSFP (*raSFP*) was eluted in five elution steps using 500 µl of denaturing elution buffer (50 mM NaH₂PO₄, 8 M urea, pH 4.5). All samples were analyzed by SDS-PAGE and western blotting. Elution aliquots were pooled and centrifuged in an Amicon tube (MWCO 3 kDa) for concentration and urea removal.

Determination of protein concentration, electrophoresis and Western blotting

Protein concentration was determined by the method of Lowry *et al.* (1951) and SDS-PAGE 14% were carried out according to Laemmli (1970) using the Mini Protean 3 system (Bio-Rad, USA). Samples were always loaded in duplicates in two gels, one for

Coomassie Brilliant Blue staining, and the other for electrotransferring to nitrocellulose membranes using a Mini Trans Blot apparatus (Bio-Rad). Membranes were incubated for 60 min at 4°C in blocking solution - Tris-buffered saline containing 5% non-fat milk (w/v) and 0.1% Tween-20 (v/v) - and further incubated with the primary antibody diluted in blocking solution overnight at 4°C. The primary antibody used was His-tag monoclonal antibody (Invitrogen), used at a concentration of 1:10,000. The membranes were washed and incubated with horseradish peroxidase-conjugated anti-mouse antibody (1:5,000) for 2 h. The chemiluminescence (ECL) was detected using X-ray films.

Binding of *raSFP* to bovine spermatozoa

The ability of spermatozoa to bind *raSFP* was accessed by incubating 1×10^7 spermatozoa with 700; 2,000; 5,000 and 7,000 ng of purified protein in a final volume of 1 ml of phosphate saline buffer (PBS) at 37°C for 5 h. As negative controls, one sperm sample was incubated without *raSFP*, and one sample was incubated with another recombinant His-tag fused protein (7,000 ng of tick calreticulin). Semen samples from two bulls were obtained by artificial vagina, and the ejaculate was washed three times in PBS to remove seminal plasma. Following incubation, samples were

washed with PBS by centrifugation at 1,000 x g for 5 min at 20°C (Dostàlovà *et al.*, 1994). To confirm the binding of *raSFP* to spermatozoa, 5×10^6 cells were lysed and protein extracts were analyzed by Western blotting as described above. The topography of *raSFP* binding pattern was studied by indirect immunofluorescence (Dostàlovà *et al.*, 1994). Briefly, 10 µl of samples (approximately 5×10^6 cells) were spread on slides, air dried and fixed in methanol. Cells were incubated for 2 h at 37°C with a 1:500 (v/v) dilution of mouse anti-His-tag antibody in PBS containing 5% BSA. The slides were washed and were incubated for another 2 h at 37°C with a 1:200 (v/v) dilution of an R-Phycoerythrin labeled goat anti-mouse IgG antibody (Sigma, USA). Slides were observed under a fluorescence microscope (Olympus, Japan). One hundred *raSFP* marked cells per slide were counted and group (four slides per group) values were statistically compared through ANOVA, followed by Tukey test assuming $P < 0.05$ as significant (Prism 6 software, GraphPad, USA).

Results

PCR amplification of aSFP cDNA from a bovine seminal vesicle library resulted in one band of approximately 400 bp by agarose gel electrophoresis analysis. The amplicon was successfully cloned into the prokaryotic expression plasmid pET-23a(+; Fig. 1).

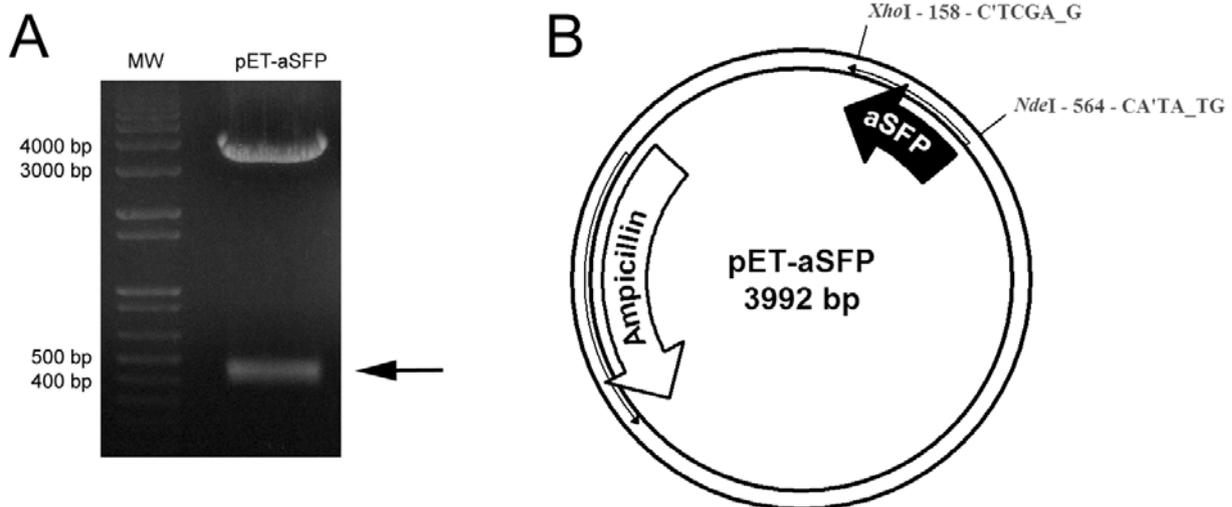


Figure 1. Construction of pET-aSFP expression vector. (A) Restriction analysis of recombinant plasmid pET-aSFP in Agarose Gel 1%. Lane MW= DNA marker; Lane 1= plasmid pET-aSFP digested with *NdeI* and *XhoI*. Arrow indicates the target gene. (B) Schematic diagram of the construct pET-asFP, showing the coding region of aSFP gene under control of T7 promoter.

Induction of *raSFP* expression was optimized after several expression conditions assays and best results were obtained with addition of 0.5 mM IPTG to the culture at 37°C for 3 h. Following sonication and centrifugation of the bacterial culture, the supernatants and inclusion bodies

were run on 12% SDS-PAGE. We obtained an expression level of approximately 30% of the total bacteria proteins (Fig. 2A). Western blotting analysis revealed that the majority of *raSFP* was in the insoluble fraction as a consequence of inclusion bodies formation (Fig. 2B and C).

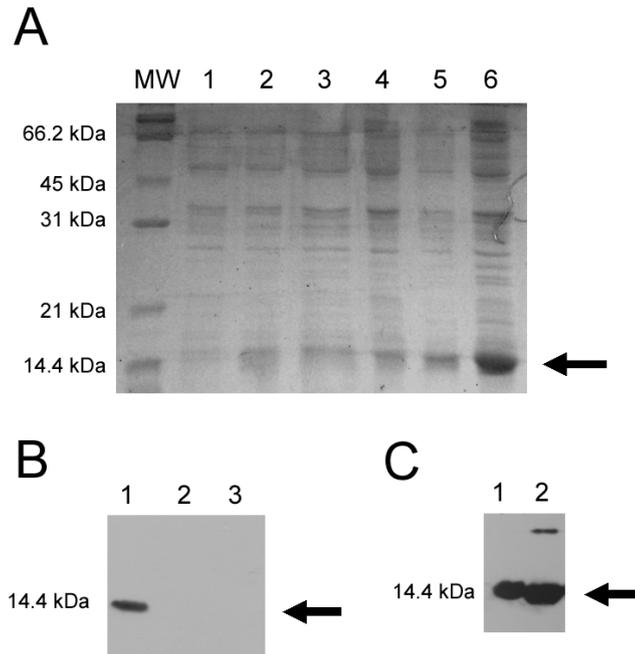


Figure 2. Analysis of *raSFP* expression. (A) Coomassie-stained SDS-PAGE of bacterial extracts. Lane MW= Molecular weight standard; Lane 1= Non induced control, 1 h; Lane 2= 1.0 mM IPTG, 1 h; Lane 3= 0.5 mM IPTG, 1 h; Lane 4= Non induced control, 3 h; Lane 5= 1.0 mM IPTG, 3 h; Lane 6= 0.5 mM IPTG, 3 h. (B) Anti-His-tag Western blot of BL21 pLysS bacterial extracts induced with 0.5 mM IPTG for 3 h. Lane 1= BL21 pLysS harboring pET-aSFP; Lane 2= BL21 pLysS harboring pET23a(+); Lane 3= BL21 pLysS without plasmid. (C) Anti-His-tag Western blot of BL21 pLysS lysates. Lane 1= supernatant; Lane 2= pellet. Arrows indicate the recombinant protein.

pET23(a)+ expression vector permits the production of N-terminal 6xHis-tagged recombinant proteins, which can be purified by one step metal chelating affinity chromatography (IMAC). Recombinant aSFP was purified using denaturing

conditions, after inclusion bodies solubilization with 8 M urea (Fig. 3A). Following protein concentration and removal of urea from solution, the SDS-PAGE profile of the pool of elution fractions showed a purity of 70% (Fig. 3B).

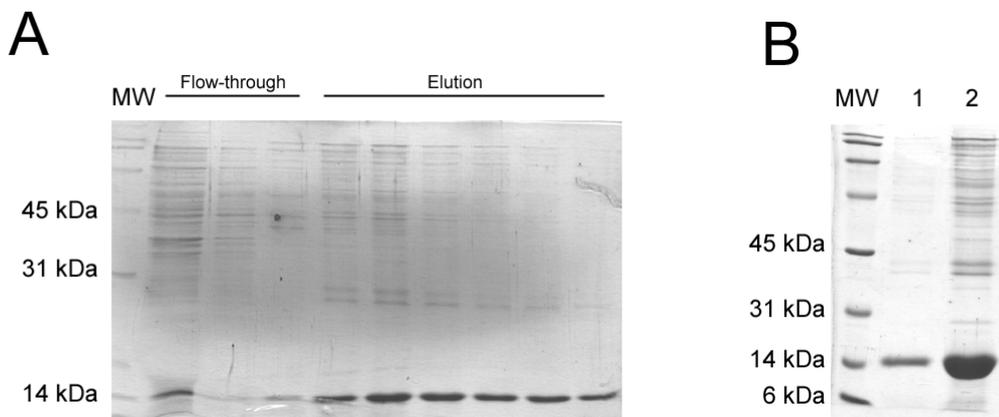


Figure 3. Purification of *raSFP* under denaturing conditions. (A) Coomassie-stained SDS-PAGE of column fractions following purification of *raSFP* on a His-Select column. (B) Coomassie-stained SDS-PAGE gel of column fractions before and after urea removal and concentration of the elution fractions. Lane 1= before concentration; Lane 2= after concentration.

Western blotting demonstrated that *raSFP* bind to ejaculated spermatozoa. Protein binding were detected in samples incubated with 700; 2,000; 5,000 and 7,000 ng of purified *raSFP* (Fig. 4). The fluorescence microscopy

confirmed the binding of *raSFP* to spermatozoa incubated with of recombinant protein (Fig. 5A), with a higher number of *raSFP*-bound spermatozoa in samples incubated with 7,000 ng of protein ($P < 0.001$, Fig. 5D). No

fluorescence signal was detected in samples incubated with 700; 2,000 ng of recombinant protein (data not shown).

Nonspecific cross reactivity was discarded by the absence of labeling on negative controls (Fig. 4, 5B and 5C).

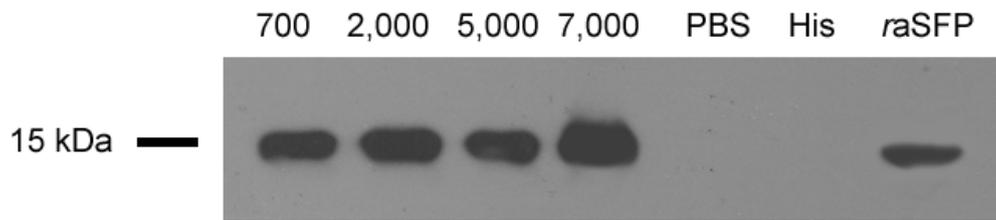


Figure 4. Binding of *raSFP* to bovine spermatozoa. Cell protein extracts of 5×10^6 spermatozoa were run in 14% SDS-PAGE, transferred to nitrocellulose membrane and incubated with different concentrations of *raSFP* from 700 to 7000 ng. Peroxidase conjugated with anti-His-tag antibody was used as a detection system. PBS= Negative control without *raSFP*; His= Negative control (incubation with 7,000 ng of his-tagged recombinant tick calreticulin); run *raSFP*= Positive control.

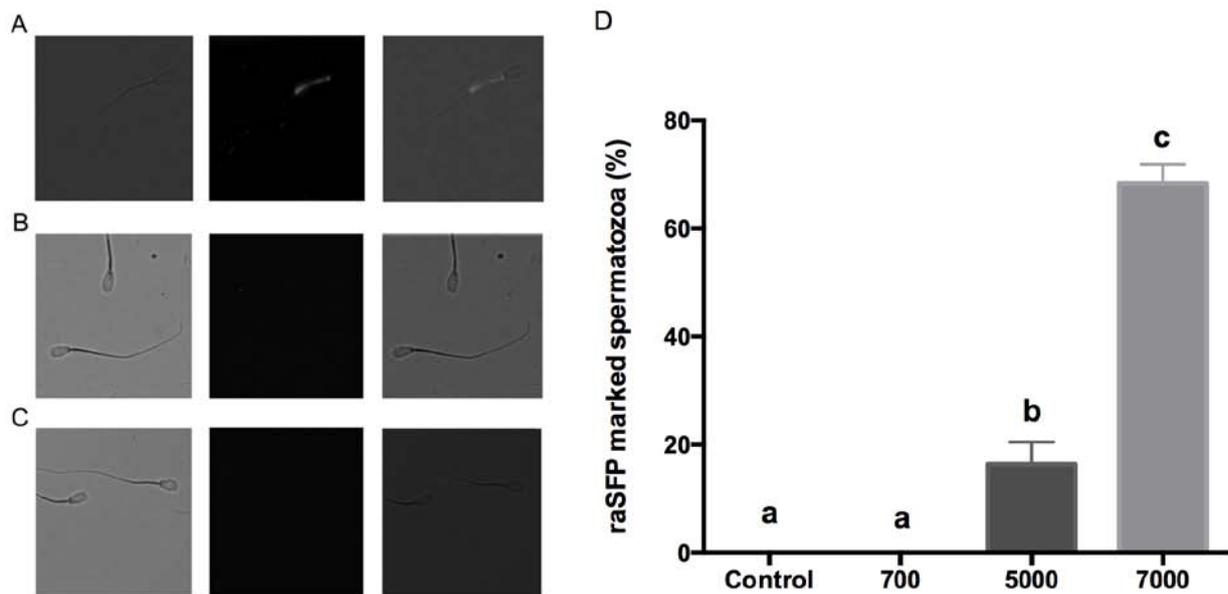


Figure 5. Representative images of immunofluorescence microscopy of bovine spermatozoa incubated with *raSFP*. Columns from left to right, microphotography under visible light, UV and merged image. A= Semen sample incubated with 7,000 ng of *raSFP*; B= Semen sample incubated in PBS; C= Semen sample incubated with His-tagged tick calreticulin. Magnification of 400X. D= Comparison of the percentage of *raSFP*-marked sperm between groups. Letters mean statistical difference ($P < 0.001$).

Discussion

Seminal plasma proteins hold important functions on sperm viability in both male and female reproductive tracts. Proteomic studies provided a broader view of the bovine seminal plasma protein content (Desnoyers *et al.*, 1994; Frazer *et al.*, 1996; Manjunath and Thérien, 2002; Jobim *et al.*, 2004; Moura *et al.*, 2007) and provided valuable information for a better understanding of their role on sperm physiology and fertilization.

Here we describe the cloning and purification of the bovine aSFP. The protein was expressed as inclusion bodies, which has many advantages including easy separation by centrifugation, protection from proteolytic degradation and lower contamination with

other *E. coli* proteins (Addona *et al.*, 2009; Meehan *et al.*, 2010). Results from Western blotting confirmed the ability of *raSFP* to attach to sperm cells after incubation with 700 ng of purified protein. The acidic Seminal Fluid Protein structure is mainly formed by a CUB domain (Romero *et al.*, 1997). Therefore, since the CUB domain may be responsible for the protein attachment to cellular membrane, we assume that *raSFP* was correctly folded after urea removal. The CUB domain is a widely occurring structural motif found almost exclusively in extracellular and plasma membrane-associated proteins (Bork and Beckmann, 1993). These proteins participate in a wide range of biological functions, including complement activation, tissue repair, cell signaling, inflammation and receptor-mediated endocytosis (Blanc *et al.*, 2007). Although the roles of the CUB domains



are yet to be completely elucidated, a number of them have been shown to be involved in oligomerization and/or recognition of substrates and binding partners (Bork and Beckmann, 1993; Blanc *et al.*, 2007).

Recombinant aSFP attached clearly to spermatozoa's mid-piece, a binding pattern similar to other spermadhesins like porcine AQN1 and AWN, which bind to the acrosome and participate in the fertilization process (Sanz *et al.*, 1992; Haase, 2005). aSFP is not related to the acrosomal function and oocyte fusion since it is believed that the protein detaches from sperm head *in utero* (Dostàlovà *et al.*, 1994). Thus, the topographical binding pattern of raSFP supports its known influence on mitochondrial activity of sperm cells. It is noteworthy to mention that spermadhesins are multifunctional proteins in other species (Topfer-Petersen *et al.*, 1998; Haase *et al.*, 2005), so different binding behaviors could be possible. To elucidate the precise function of aSFP on motility preservation of frozen semen, it is important to investigate the binding mechanism to sperm membrane and which signaling pathways aSFP might be interacting with. Differently from other spermadhesins, native and recombinant aSFP do not possess mannose binding properties (Ekhlasi-Hundrieser *et al.*, 2008), so a protein-protein interaction is more likely. However, no specific protein receptor for aSFP in sperm membrane has been described so far.

Proteomic studies suggested aSFP as a marker for high semen freezability (Jobim *et al.*, 2004), however its potential role on sperm preservation is yet to be determined. The influence of aSFP in bovine semen was demonstrated by (Schöneck *et al.*, 1996). The addition of a high concentration of bovine SP purified aSFP in semen, decreased sperm motility and mitochondrial activity. Since aSFP is secreted at the epididymis, a natural sperm reservoir, these effects lead to a lower production of reactive oxygen species (ROS) by stored sperm cells. In addition, aSFP showed a remarkable antioxidant behavior due to its redox equilibrium exhibited by the cysteine residues Cys54 and Cys75 (Calvete and Sanz, 2007). These Cys residues are responsible for aSFP alternation between oxidized and reduced states, acting as ROS scavenger.

As a result, bulls with higher semen freezability may ejaculate spermatozoa with lower oxidative damage on membrane lipids and proteins. These effects are an important strategy for semen storage in the epididymis, resulting in a better preservation of sperm motility and viability (Schöneck *et al.*, 1996). Thus, the role of aSFP on semen freezability might be associated with the quality of the stored sperm and with a minor effect after ejaculation. Egg yolk-based semen extenders present a marked antioxidant activity (Bustamante-Filho *et al.*, 2009), so aSFP antioxidant protection may only be beneficial before semen dilution.

In this article we report, for the first time, the recombinant expression and purification of aSFP. The

plasmid (pET23a) used in the expression study was successfully constructed by the ligation of the mRNA aSFP cDNA, resulting in N-terminal tagging with 6xHis. The expression of the raSFP in bacterial system (*E. coli* BL21 pLysS) and further purification was successfully accomplished and may help us understand the mechanism by which this protein modulates sperm metabolism and cryopreservation success in further studies.

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