

S100A7 in the Fallopian tube: a comparative study

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Summary

The oviduct is a dynamic organ in which final gamete maturation, fertilization and early embryo development take place. It is considered to be a sterile site; however the mechanism for sterility maintenance is still unknown. S100A7 is an anti-microbial peptide that has been reported in human reproductive tissues such as prostate, testicle, ovary, normal cervical epithelium and sperm. The current work reports the presence of S100A7 in the Fallopian tube and its localization at the apical surface of epithelial cells. For comparison, porcine S100A7 was used for antibody development and search for peptide in reproductive tissues. Although present in boar seminal vesicles and seminal plasma, S100A7 was not detected on female porcine organs. Also, in contrast with the human protein, porcine S100A7 did not show anti-microbial activity under the conditions tested. Phylogenetic analyses showed high divergence of porcine S100A7 from human, primate, bovine, ovine and equine sequences, being the murine sequence at a most distant branch. The differences in sequence homology, *Escherichia coli*-cidal activity, detectable presence and localization of S100A7 from human and pig, suggest that there are possible different functions in each organism.

Keywords: Fallopian tube, Oviduct sperm, S100A7, Seminal plasma

Introduction

In order for mammalian fertilization to proceed sperm must enter the female reproductive tract, where the selection processes submit them to strict controls (Hunter, 2005; Suárez, 2007). The subpopulation of sperm that reach the oviduct can bind to the epithelial cells of the isthmus, as occurs in several mammals such as human, sheep, cow, horse, pig and mouse (Hunter, 1981; Flechon & Hunter, 1981; Hunter & Nichol, 1983; Hunter & Wilmut, 1984; Demott & Suárez, 1992; Thomas *et al.*, 1994; Pacey *et al.*, 1995; Suárez, 1998);

such bound sperm show decreased movement and prolonged survival (Fazeli *et al.*, 2003; Suárez, 2008). It has been demonstrated that oviductal proteins can modulate those sperm functions (Avilés *et al.*, 2010; Holt & Fazeli, 2010; Hung & Suárez, 2010; Killian, 2011). Oviductal annexin A2 has been involved in the attachment of sperm to oviductal epithelium in pig (Teijeiro *et al.*, 2009) and cow (Ignatz *et al.*, 2007) and oviductal porcine deleted in malignant brain tumour 1 (DMBT1) has been related to sperm selection (Teijeiro & Marini, 2012a; Teijeiro *et al.*, 2008, 2011, 2012).

However, the female genital tract is a complex biosystem that is constantly exposed to a multitude of bacterial, fungal and viral agents (Mildner *et al.*, 2010). Importantly, anti-microbial peptides (AMPs) are small proteins produced by epithelial surfaces and inflammatory cells that have broad-spectrum anti-microbial and immunomodulatory activities and are present in a number of sites throughout the female reproductive tract (Frew & Stock, 2011). S100A7 (Psoriasin), a protein that belongs to a multigenic family of calcium-modulated proteins of the EF-hand type, has been identified as an AMP with strong

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activity against *Escherichia coli* (Gläser *et al.*, 2005). Moreover, S100A7 has been proposed as the major *Escherichia coli*-cidal factor in the female genital tract, and is expressed in vulva, vagina and ectocervical epithelia (Mildner *et al.*, 2010). In addition, expression of S100A7 has been found in reproductive tissues such as prostate, testicle, ovary and normal cervical epithelium (Shadeo *et al.*, 2007). The present authors have identified S100A7 in human and boar sperm and have demonstrated the interaction of porcine sperm S100A7 with porcine oviductal DMBT1 (Teijeiro & Marini, 2012b).

As AMPs are likely to have important functions in the female reproductive tract, and the expression of S100A7, an AMP, was not evaluated in Fallopian tubes, which are considered to be 'sterile sites' (Quayle, 2002), it was decided to search for the presence of this protein in the human oviduct.

Comparatively, the presence of S100A7 in porcine reproductive tissues was also evaluated. Because the *Escherichia coli*-cidal activity of S100A7 has been proposed in mammals other than human, such as cows (Regenhard *et al.*, 2009), anti-microbial activity and further sequence homology analysis of porcine S100A7 were performed.

Materials and methods

Chemicals

Chemicals were obtained from Sigma-Aldrich (Buenos Aires, Argentina), unless otherwise stated. The following primary antibodies were used: polyclonal anti-human S100A7 purified by affinity to His-6-S100A7 immobilized on nitrocellulose membranes (Teijeiro & Marini, 2012b) (dilution 1:1000 for western blot and 1:100 for immunofluorescence) and polyclonal anti-porcine S100A7 (dilution 1:1000). The following secondary antibodies were used: horseradish peroxidase-conjugated anti-rabbit IgG (dilution 1:10,000) (GE Healthcare, Buenos Aires, Argentina); Cy3-conjugated anti-rabbit immunoglobulin (dilution 1:1000) (Chemicon International Inc., Temecula, USA).

Sample collection

Boar seminal plasma was collected from adult fertile boars from different breeds by the glove-handed method. The sperm-rich fraction was separated from seminal plasma by centrifugation. Porcine tongues, testicles, epididymis, seminal vesicles, ovaries, oviducts and uteri were obtained from a local abattoir and transported immediately to the laboratory in ice-cold phosphate-buffered saline (PBS). Each tissue was dissected and cut into pieces 1 × 1 cm.

Then, each fragment was resuspended in buffer that contained 7 M urea, 2 M thiourea, 4% (w/v) CHAPS, 1% NP40 and dithiothreitol (DTT) 100 mM, and disaggregated by forceps and scissors. The tissue was homogenized in a Potter homogenizer and then sonicated and centrifuged 15 min at 4 °C at 16,000 g. The supernatant was collected and stored at -80 °C until use. Human seminal plasma and Fallopian tubes were gifted by Dr Sergio Ghersevich from the Clinical Biochemistry Area, Facultad de Ciencias Bioquímicas y Farmacéuticas, Universidad Nacional de Rosario, Argentina. For human samples, study protocols were approved by the Institutional Bioethical Board of Facultad de Ciencias Bioquímicas y Farmacéuticas, Universidad Nacional de Rosario, Argentina and written consent was obtained from all donors. Human oviductal tissue was obtained from premenopausal women ($n = 6$, average age: 42.9 years) with no clinical history of infection or neoplastic diseases, scheduled for hysterectomies as a result of uterine fibromyomas or hypermenorrhea (Hospital Provincial del Centenario, Rosario, Argentina). Samples for western blot were processed in accordance with Zumoffen *et al.* (2013), briefly, fragments of the oviducts were opened longitudinally and the epithelial layer of cells was carefully scraped out with a scalpel and homogenized in homogenization buffer that consisted of 50 mM Tris-HCl (pH 7.4), 0.1 mM EDTA, 0.1 mM EGTA, 0.1 M phenylmethanesulfonyl fluoride (PMSF), 2 µg/ml Aprotinin and 0.1% v/v 2-mercaptoethanol, in an ice bath. The oviductal cells homogenates were then centrifuged at 15,000 g for 30 min at 4 °C, and the supernatants stored at -20 °C until further use. Fallopian tube samples for immunofluorescence were provided as paraffin blocks.

Immunofluorescence

Human Fallopian tube tissues were sectioned, deparaffinized in xylene, and then rehydrated through graded dilutions of ethanol (100, 95, 70 and 35%), followed by two washes in Tris-buffered saline (TBS) (25 mM Tris-HCl pH 7.5, 150 mM NaCl). Slides were gently rinsed twice with TBS, blocked with 2% BSA, 0.2% Triton X-100 in TBS for 1 h and treated with the primary antibodies overnight at 4 °C. After rinsing twice with TBS, the slides were treated with Cy3-conjugated anti-rabbit immunoglobulin (1:2,000) for 1h. After rinsing twice with TBS, slides were covered with 4',6-diamidino-2-phenylindole (DAPI) for 30 min, then were washed with TBS, covered with 0.22 M 1,4-diazabicyclo[2,2,2]octane dissolved in glycerol:TBS (9:1) and cover slips were applied. The preparations were examined under a confocal microscope (Nikon Model Eclipse TE-2000-E2, Melville, USA).

Subcloning and expression of porcine S100A7, and development of anti-porcine S100A7 antibodies (anti-His6-pS100A7)

The coding sequence for porcine S100A7 (pS100A7) was cloned into pBluescript II by Gorodkin *et al.* (2007) and was a gift from Dr Claus Jørgensen (Faculty of Life Sciences, University of Copenhagen, Denmark). This sequence was subcloned into pET-tev vector as histidine fusion and the recombinant protein (His6-pS100A7) was expressed in *E. coli* BL21. Protein expression was induced with isopropyl-1-thio- β -D-galactopyranoside (IPTG) for 20 h and purification was performed with Ni-NTA agarose (QIAGEN, Hilden, Germany) in accordance with the manufacturer's instructions. His6-pS100A7 was eluted five times sequentially with 250 mM imidazole; 0.5 ml fractions were collected and subsequently tested for purity by silver-stained SDS-PAGE (Rabilloud *et al.*, 1994). Fractions that contained His6-pS100A7 were pooled and concentrated by ultrafiltration using Ultracel-3 K (Amicon® Ultra, Millipore, Massachusetts, USA) and re-buffered into PBS (130 mM NaCl, 2.68 mM KCl, 6.46 mM Na₂HPO₄, 1.47 mM KH₂PO₄), pH 7.4.

Purified recombinant protein was used for anti-serum development in rabbit as described by Pérez *et al.* (2006). Polyclonal anti-human S100A7 antibodies (anti-hS100A7) were developed in rabbit using recombinant GST-hS100A7 expressed and purified as described by Teijeiro & Marini (2012a).

Western blots

Protein extracts obtained as mentioned above were supplemented with loading buffer (25 mM Tris-HCl, pH 6.8, 30% glycerol, 4% SDS) boiled for 5 min, used for 15% SDS-PAGE and then proteins were transferred to nitrocellulose membranes (GE Healthcare, Buenos Aires, Argentina). Non-specific binding sites were blocked by incubation with 5% dried non-fat milk in TBS (25 mM Tris-HCl, pH 7.4, 150 mM NaCl). Membranes were treated 1 h with the corresponding antibodies. After washing (three times for 10 min each), the blots were incubated with peroxidase-conjugated anti-rabbit IgG for 1 h and washed again. Labelled proteins were revealed using enhanced chemiluminescence detection with ECL Kit (GE Healthcare, Buenos Aires, Argentina).

Microdilution susceptibility assay

The microdilution susceptibility assay was performed as described by Regenhard *et al.* (2009) with minor modifications. Only *Escherichia coli* strain ATCC 35218 was used for the microdilution assay. In brief, cells were grown for 2 h in Luria-Bertali (LB) medium at 37°C. Then the bacterial culture was adjusted to 10⁴

colony-forming units (CFU)/ml in 10 mM sodium phosphate buffer (pH 7.4).

Aliquots (100 μ l) of the microbial suspension were mixed with 10 μ l of His6-pS100A7 solution (range of final concentration from 50–150 μ g/l) and incubated at 37°C. After 3 h, the total volume of test suspensions was plated onto LB-Agar. After overnight incubation at 37°C, CFU were counted. Results are given as CFU/ml. Two-way analysis of variance (ANOVA) was applied and individual means were further tested by least significant difference (LSD) test.

Radial diffusion assay

A radial diffusion assay was performed as in Lehrer *et al.* (1991) with modifications. Bacteria were grown to mid-log phase in LB medium and plated into Petri dishes (10 ml). After plating, a 5 μ l aliquot of each protein solution was pipetted into wells formed with a biopsy punch (diameter: 3 mm), and plates were incubated overnight at 37°C. Ampicillin and kanamycin were used as positive growth inhibition controls.

Phylogeny and sequence analysis of S100A7 proteins in mammals

In order to broaden the study to other mammals, the presence of S100A7 proteins from other mammalian species was sought. The proteins that were manually annotated and reviewed were used for comparison in their annotated form. The proteins that were automatically annotated and not reviewed were compared with the previous ones using a BLAST (Basic Local Alignment Search Tool) (Altschul *et al.*, 1997) program to discard redundant data. Accession numbers for S100A7 proteins used in this analysis were: gi|190668 (*Homo sapiens*), gi|114559708 (*Pan troglodytes*), gi|27807077 (*Bos taurus*), gi|110224792 (*Equus caballus*), gi|46397569 (*Mus musculus*), gi|426216655 (*Ovis aries*), EW648856.2 (*Sus scrofa*) and gi|350583366 (*Sus scrofa*). Phylogenetic analyses were carried out using <http://www.phylogeny.fr>. Multiple sequence alignments were performed using the MUSCLE algorithm. Curation was performed using G-blocks with non-stringent selection parameters. A phylogenetic tree was reconstructed using maximum of likelihood (PhyML) method. Bootstrap values were estimated with 100 replications.

Molecular weight of proteins was estimated from the sequences using bioinformatics tools from <http://expasy.org/>

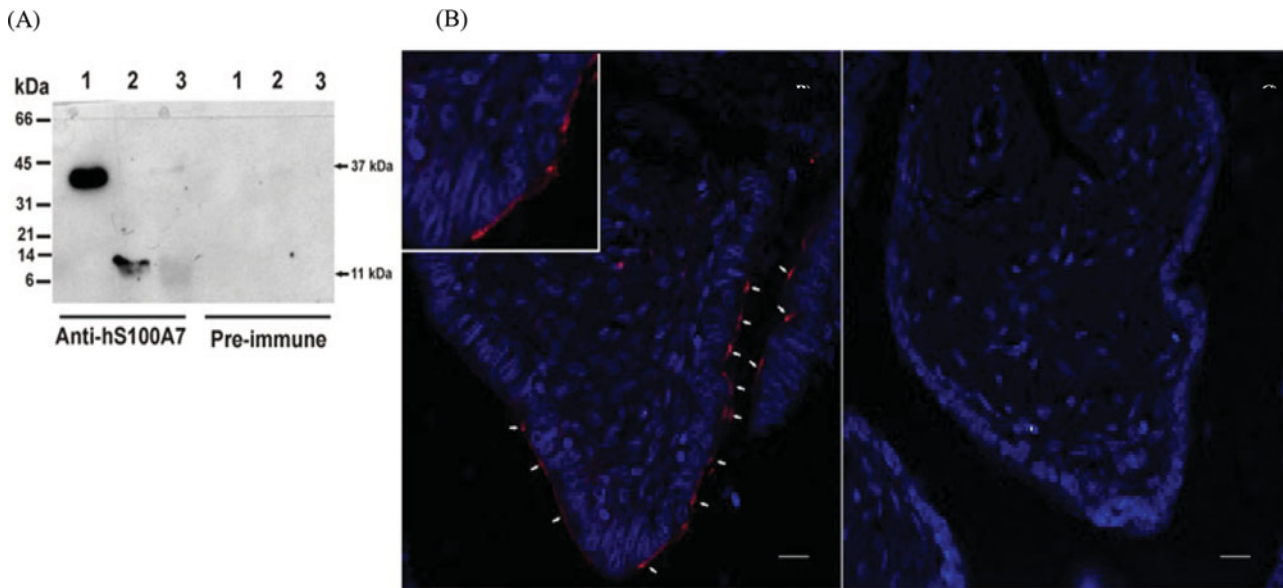


Figure 1 Human S100A7 expression in Fallopian tube and seminal plasma. (A) Western blot analyses of Fallopian tube tissue extracts and seminal plasma. Line 1: 2 μ g of GST-hS100A7; line 2: Fallopian tube tissue extract (protein content 50 μ g); line 3: seminal plasma (protein content 30 μ g). Data are representative of six Fallopian tubes and five seminal plasma samples. (B) Immunofluorescence showing hS100A7 in the apical cytoplasm of epithelial cells (white arrows). Inset shows $\times 2.5$ magnification. (C) Control was performed with pre-immune serum. Nuclei were stained with DAPI. Bar indicates 20 μ m.

Results

Presence of S100A7 in Fallopian tubes

To evaluate the presence of S100A7 in Fallopian tubes (human S100A7, hS100A7), tissue protein extracts were subjected to western blot assays. Antibodies generated against recombinant GST-hS100A7 were purified by affinity to recombinant His6-hS100A7 and used for this assay (Teijeiro & Marini, 2012b). Fig. 1A shows the presence of hS100A7 in protein extracts from Fallopian tubes and in seminal plasma. Immunofluorescence analysis showed that hS100A7 is located at the apical cytoplasm of Fallopian epithelial cells (Fig. 1B, white arrows, left panel). Negative controls were developed with pre-immune serum (Fig. 1C, right panel).

Presence of S100A7 in porcine tissues

Porcine S100A7 was initially cloned from tongue RNA (Gorodkin *et al.*, 2007). To corroborate the protein expression in its original tissue resource and to evaluate the quality of the antiserum prepared using His6-pS100A7, protein extracts from this organ were subjected to western blot. As shown in Fig. 2A, different amounts of a band of 11 kDa that corresponded to pS100A7 were observed for different quantities of tongue extracts revealed with anti-His6-pS100A7. Pre-immune serum showed no signal. Having checked the quality of anti-His6-pS100A10 antiserum, the presence of pS100A7 in different reproductive tissues was

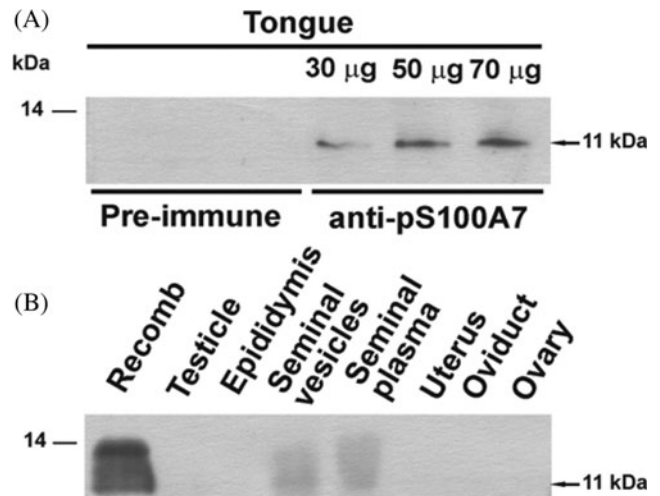


Figure 2 Presence of S100A7 in porcine tissues. (A) Expression of pS100A7 in tongue. Different amounts of tissue extracts were assayed by western blot showing increased amounts of pS100A7. Control was performed with pre-immune serum. (B) Analysis of expression of pS100A7 in porcine reproductive tissues by western blot. Only seminal vesicles and seminal plasma showed signal. Tissue extracts containing 50 μ g of protein were used. Recomb: recombinant pS100A7 (2 μ g).

evaluated by western blot. From all the assayed reproductive tissues, only the seminal vesicles showed the presence of pS100A7. Seminal plasma also showed

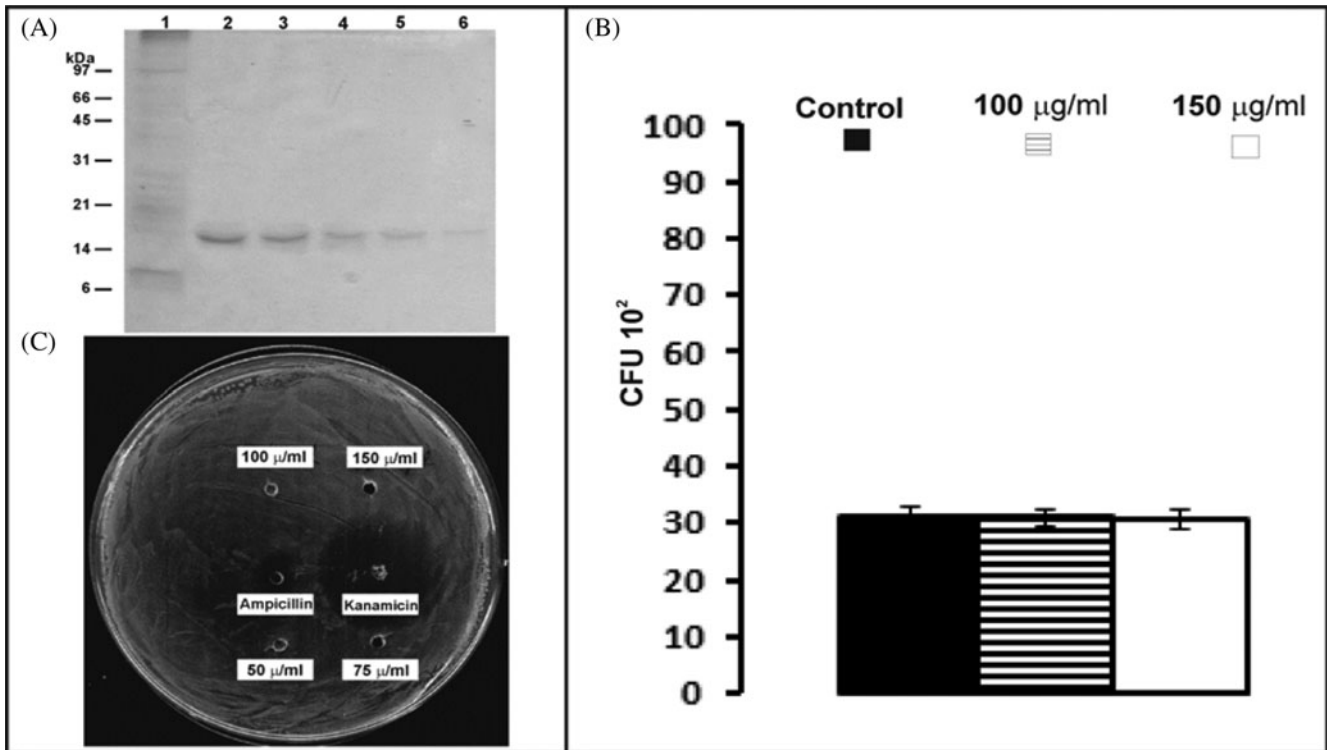


Figure 3 Analyses of His6-pS100A7 anti-microbial activity. (A) Purification of recombinant pS100A7: 10 microliter samples from purified aliquots sequentially eluted of His6-pS100A7 were analysed by silver-stained SDS-PAGE (lines 2–6). Line 1: 10 µg *E. coli* BL21 IPTG induced culture extract. (B) *E. coli* ATCC 35218 was treated with purified His6-pS100A7. Only two of the highest concentrations used are shown. Microdilution susceptibility assay shows no differences in CFU counting between treatments and control. (C) Radial diffusion assay shows no clear zone surrounding the wells loaded with different concentrations of His6-pS100A7. Ampicillin and kanamycin were used as positive controls.

signal. S100A7 was not detected on porcine female organs under these conditions.

His6-pS100A7 anti-microbial activity analyses

Previous reports indicate that cow S100A7 and human S100A7 have anti-microbial activity, preferentially against *E. coli*. Sequentially eluted His6-pS100A7 fractions were purified with the efficacy illustrated in Fig. 3A, pooled and re-buffered with PBS. Purified His6-pS100A7 was then used for anti-microbial activity assays. Testing the effect on the same *E. coli* strain, ATCC 35218, by microdilution susceptibility and radial diffusion assays, no anti-microbial activity for recombinant pS100A7 could be detected. There was no significant difference between CFU in *E. coli* treated with His6-pS100A7 and controls (Fig. 3B). Also, complementary radial diffusion assay showed no clear zones in wells that contained different concentrations of recombinant pS100A7, which indicated no inhibition of *E. coli* growth by pS100A7, in comparison with well known anti-microbial compounds (Fig. 3C).

Phylogeny analysis of S100A7 proteins in mammals

A phylogenetic tree was reconstructed using all the sequences indicated in the Materials and methods section. The phylogenetic reconstruction and the multiple alignment data showed that the amino acid sequence of *Mus musculus* (gi | 46397569) and *Sus scrofa* (EW648856.2) S100A7 are highly divergent from the other sequences (Fig. 4A, B). As would be expected from patterns of mammalian evolution, primate and ruminant S100A7 cluster together. The phylogenetic tree reconstructed using the PhyML method suggests that the equine homologue is more closely related to primate sequences, while porcine and murine proteins form different branches.

Analysis of molecular weights, calculated using bioinformatics tools, showed that the weight for the automatically derived porcine S100A7 sequence (gi | 350583366) was 17621.97 Da, while that for tongue cDNA cloned porcine S100A7 (EW648856.2) was 11653.99 Da (Fig. 4C, in bold). The latter was confirmed to be present in tongue tissue by western blot in the current study (Fig. 2A). It should be noted that the amino acid at position 99 of the polypeptide derived

in the oviductal epithelium (Fig. 2B). The presence of hS100A7 was detected in human testicles and ovaries (Shadeo *et al.*, 2007) but pS100A7 was not detected in the same tissues from pigs. Thus, S100A7 seems to exhibit a conserved role in the tongue that is not observed between human and porcine reproductive tissues.

Anti-microbial activity for His6-pS100A7 was not detected either by microdilution susceptibility assay or by radial diffusion assay (Fig. 3B, C); however, this activity cannot be ruled out.

As expected from mammalian pattern evolution, the current phylogenetic reconstruction shows that primates and ruminants cluster together (Fig. 4). Although other primate S100A7 sequences are available, for example *Gorilla gorilla* (gi|426331624), they show extremely high homology and thus chimpanzee was chosen as representative of the group. Equine S100A7 is more related to S100A7 from primates, but no information could be found about its functions or localizations. In agreement with Wolf *et al.* (2006), the current phylogenetic reconstruction suggests that murine S100A7 may be the ancestral gene for human S100A7 and, moreover, for the rest of the species analysed. Porcine S100A7 forms a different branch from that of most species, and this might be related to the lack of the anti-microbial activity shown, in contrast to the bovine and human proteins. Mutation of the carboxyterminal region of human S100A7 slightly reduces psoriasis's anti-bacterial activity (Lee & Eckert, 2007). In this regard, the lack of homology between the porcine and human proteins at the carboxyterminal end might be responsible for the lack of anti-microbial activity of pS100A7, and indicates that innate immunity may not be the main function of S100A7 in porcine reproductive tracts. Bioinformatic tools derived the molecular weight for the automatically annotated porcine protein (gi|350583366) (Fig. 4) as 17621.97 Da, while that of S100A7 cloned from pig tongues was 11653.99 Da. The tongue-cloned molecular weight is in agreement with the weight of the actual protein observed by western blot. The apparent molecular mass corresponds to the sequence obtained from tongue cDNA instead of the one deduced from the DNA sequence (gi|350583366) (Fig. 4C).

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