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## Author Contribution

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Yan

“revised”

**1 Decreasing of S100A4 in bovine endometritis in vivo and in vitro**

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20 **Abstract**

21 Endometritis is a prevalent reproductive disease in dairy cows, and is a  
22 superficial inflammation of the endometrium. S100 calcium-binding protein A4  
23 (S100A4) is suggested to be implicated in the progression of inflammation. However,  
24 to our knowledge, no study has reported the changes of S100A4 during bovine  
25 endometritis. The objective of this study was to investigate S100A4 gene expression  
26 and protein levels in the uterus with endometritis in dairy cows. Vaginal mucus  
27 samples were collected for diagnosis of the severity degree of endometritis and the  
28 detection of S100A4 protein content. Blood samples and endometrial biopsies were  
29 collected and divided into the control (CN), mild endometritis (M), and severe  
30 endometritis (S) groups according to the characteristics of the vaginal mucus type.  
31 The isolated bovine endometrial epithelial cells (BEECs) were challenged with *E. coli*  
32 ( $2 \times 10^6$  CFU/mL,  $2 \times 10^7$  CFU/mL) or lipopolysaccharide (LPS, 3 and 10  $\mu$ g/mL) as an  
33 inflammatory model. RT-qPCR was used to detect the gene expression levels of  
34 S100A4 and cytokines, including interleukin-1 $\beta$  (IL-1 $\beta$ ), interleukin-6 (IL-6),  
35 interleukin-10 (IL-10), and tumour necrosis factor-alpha (TNF- $\alpha$ ), in tissues or cells.  
36 Enzyme-linked immunosorbent assay (ELISA) was used for S100A4 protein level  
37 detection in tissues, cells, cell supernatant, vaginal mucus, and serum samples. The  
38 results showed that S100A4 gene and protein levels decreased in bovine endometrium  
39 with endometritis and in *E. coli*- or LPS-stimulated BEECs. We failed to detect  
40 S100A4 in the cell supernatant, vaginal mucus, and serum samples. This study  
41 suggested that S100A4 is a pathogenesis-related protein of endometritis, and

42 decreased expression of S100A4 may pave the way for the development of  
43 endometritis in dairy cows.

44 **Key words:** S100A4, endometritis, dairy cow

## 45 **1. Introduction**

46 As a result of milk accounting for a large proportion of the human diet, the health  
47 conditions of dairy cows have attracted great attention. Reproductive disease,  
48 especially endometritis, is one of the most important diseases compromising animal  
49 welfare and causing increased economic loss in the dairy industry [1]. Endometritis is  
50 highly prevalent in dairy cows and can result in other kinds of reproductive diseases,  
51 including pyometra, anovulation and pregnancy loss, as well as reduce productive  
52 performance [2, 3].

53 S100 calcium-binding proteins are widely reported to be crucial to many diseases  
54 and are considered potential biomarkers in the diagnosis and treatment of several  
55 diseases [4]. S100A4 is an S100 protein family member that functions in both  
56 intracellular and extracellular pathways[5]. Existing studies have shown that S100A4  
57 protein is expressed in immune cells, such as monocytes, macrophages, and  
58 polymorphonuclear granulocytes, and is also expressed at low levels in normal tissues  
59 [6, 7]. S100A4 plays important roles in a range of biological functions, including cell  
60 adhesion, movement, invasion, and metastasis [8]. Numerous studies on S100A4 have  
61 focused on tumours and cancer [5]. S100A4 has also been reported to be involved in  
62 inflammation progression. For example, S100A4 can aggravate the clinical symptoms  
63 of colitis in mice [9], and it may be a potent trigger of inflammatory processes and

64 induce the release of cytokines [10]. It has also been reported that S100A4 promoted  
65 the progression of endometritis, while deficiency of S100A4 reduced the mRNA  
66 expression levels of IL-1 $\beta$  and TNF- $\alpha$  in the endometrium of mice [11]. However, the  
67 roles of S100A4 in bovine endometritis have not been reported. To investigate the  
68 roles of S100A4 in endometritis of dairy cows, we determined the gene and protein  
69 expression of S100A4 in endometrial tissues, vaginal mucus, serum, *E.*  
70 *coli*/LPS-stimulated endometrial epithelial cells, and cell supernatant.

## 71 **2. Materials and methods**

### 72 *2.1. Animals and diagnostic samples*

73 We randomly selected twenty-two Holstein-Friesian cows, aged 3 to 5 years and  
74 2 to 3 parities on 40 to 60 days postpartum without systemic symptoms. Vaginal  
75 discharge samples were collected using a draw-tube. All cows were divided into three  
76 groups based on discharge detection: control group (CN, n=6) characterized by clear  
77 vaginal mucus, mild endometritis group (M, n=8) characterized by less than 50%  
78 white or off-white mucopurulent material in vaginal discharge, and severe  
79 endometritis group (S, n=8) characterized by more than 50% white or yellow purulent  
80 material in vaginal discharge [12]. Endometrial biopsy samples were obtained using  
81 sterile uterine biopsy punch forceps from the right side of the uterine horn. The tissue  
82 samples were divided into two parts: one was immediately placed into liquid nitrogen  
83 for total protein and RNA extraction, and another was placed into 10% neutral  
84 formalin for histopathology using routine haematoxylin and eosin (HE) staining.

85 Bovine endometrial epithelial cells (BEECs) were isolated from a healthy uterus  
86 of a six-month old dairy cow. Briefly, a healthy uterus was brought to the laboratory  
87 in a sterile PBS (pH 7.2) containing penicillin (100 µg/mL)  
88 and streptomycin (100 U/mL). Endometrium from the uterine horn was cut off into 2–  
89 3 cm long pieces, washed in PBS (pH 7.2) twice, and then uterine tissue was digested  
90 with 1% collagenase I (Sigma, USA) diluted in DMEM/F12 (HyClone, USA) for 6 h.  
91 The digested endometrium was scraped using a sterile cell scraper, and scraped  
92 materials were collected and washed in PBS (pH 7.2). Then, the collected materials  
93 were centrifuged at  $100\times g$  for 5 min to collect the cell suspension. Trypan Blue stain  
94 was used to estimate cell viability. Cells were cultured in DMEM/F12 with 10% foetal  
95 bovine serum (Gibco, Australia), penicillin (100 µg/mL), and streptomycin (100 U/mL)  
96 at 37 °C with 5% CO<sub>2</sub> and 95% sterile air when the viability >95%. The medium was  
97 changed every 3 days until the cells reached approximately 90% confluence.

98 Considering that the cultures were inevitably mixed with some stromal cells, we  
99 removed stromal cells according to the different sensitivities of epithelial cells and  
100 stromal cells to trypsin [13]. We conducted time-different digestion thrice to obtain  
101 purified epithelial cells. We used the expression of epithelial-specific cytokeratin to  
102 estimate the purification of epithelial cells using a mouse monoclonal antibody  
103 (Abcam, ab49779) and goat anti-mouse IgG-FITC (Abcam, ab6785).

104 Cells growing in logarithmic phase were inoculated into a six-well plate at a  
105 concentration of  $2\times 10^5$  cells/well, and incubated for 24 h as described above. They  
106 were then treated with different concentrations of *E. coli* ( $2\times 10^6$  CFU/mL,  $2\times 10^7$

107 CFU/mL) for 2 h or LPS (O111:B4, Sigma) at 0, 3, or 10  $\mu\text{g/mL}$  for 24 h. After  
108 stimulation, the cells and supernatant were collected for gene or protein analysis of  
109 S100A4. Three biological replicates and three technological replicates were included.

## 110 2.2. Diagnostic assays

111 Total RNA was extracted from BEECs or endometrial biopsy tissues using  
112 RNAiso Plus (Takara, Japan), and 600 ng of RNA was used for reverse transcription  
113 by the PrimeScript RT reagent Kit with gDNA Eraser (Takara, Japan). RT-qPCR was  
114 performed using SYBR Premix Ex Taq  $\square$  (Takara, Japan) on an iQ5 thermal cycler  
115 (Bio-Rad, USA). The fold change in mRNA expression relative to  $\beta$ -actin was  
116 calculated using the  $2^{-\Delta\Delta C_t}$  method. The involved primers are detailed in Table 1.

117 A commercial kit (Transgen, China) was used for total protein extraction in  
118 endometrial biopsies and cells. Briefly, grounding tissues (ground with liquid nitrogen  
119 using a mortar) and cells were treated with lysis solution to release total protein. Then,  
120 the samples were centrifuged at  $14,000\times g$  for 10 min at  $4^\circ\text{C}$  to collect total protein.  
121 The concentration of total protein was measured using a BCA Protein Assay Kit  
122 (Takara, Japan).

123 We used a commercial ELISA kit for human beings (CUSABIO, China) to  
124 determine the S100A4 level in total protein samples from tissues and cells, cell  
125 supernatant, vaginal mucus, and serum samples. The detection range of the S100A4  
126 contents using this ELISA kit is 0.9 ng/mL to 60 ng/mL. MULTISKAN MK3  
127 (Thermo, USA) was used to read the plate. Serum was isolated from blood samples by  
128 centrifugation at  $3000\times g$  for 10 min. Pre-treatment of vaginal mucus was performed



129 before ELISA detection. The mucus was diluted with sterile PBS containing  
130 protease inhibitor, then centrifuged at  $12,000\times g$  for 10 min. A BCA Protein Assay  
131 Kit (Takara, Japan) was used to measure the concentration of total protein in mucus.

### 132 2.3. Statistical analysis

133 Statistical data analysis was carried out with analysis of variance built in SPSS  
134 19 (Chicago, IL, USA). All data are presented as means  $\pm$  SD. One-way ANOVA  
135 analysis was used to compare the results among groups. Statistical significance was  
136 considered at  $P<0.05$ . Differences among all groups at each index were performed.  
137 Totally different letters indicated significant difference between two group ( $P<0.05$ ).

## 138 3. Results

139 As shown in Fig. 1, the healthy endometrium tissues had a complete epithelium  
140 composed of normal epithelial cells; in mild endometritis tissues, the epithelial cells  
141 were arranged irregularly in the endometrium with prominent lymphocyte-rich  
142 infiltration and low-grade plasmocyte infiltration; the endometrium tissues with  
143 severe endometritis showed lymphocyte migration, haemorrhage, and epithelial  
144 necrosis. The histopathological features of the three groups were consistent with  
145 vaginal mucus classification. The gene and protein expression of S100A4 in the  
146 classified tissue samples was detected, and the results showed that both the gene  
147 expression level and protein content of S100A4 were significantly decreased in  
148 endometrial tissues with mild or severe endometritis (Fig. 2) and negatively correlated  
149 with the severity of endometritis.

150 The results of the in vitro experiment indicated that, compared with normal  
151 BEECs, TNF- $\alpha$ , IL-1 $\beta$ , IL-6, and IL-10 were upregulated significantly when BEECs  
152 were stimulated with *E. coli* ( $2 \times 10^6$  CFU/mL,  $2 \times 10^7$  CFU/mL), with the exception of  
153 IL-6 levels after  $2 \times 10^6$  CFU/mL *E. coli* stimulation (Fig. 3A-D). Likewise, the mRNA  
154 levels of TNF- $\alpha$ , IL-1 $\beta$ , IL-6 and IL-10 increased significantly when BEECs were  
155 treated with 3 or 10  $\mu$ g/mL LPS (Fig 3E-H). Nevertheless, S100A4 gene levels  
156 significantly decreased in BEECs treated with *E. coli* ( $2 \times 10^6$  CFU/mL,  $2 \times 10^7$   
157 CFU/mL) or LPS (3, 10  $\mu$ g/mL), and protein levels also significantly decreased in  
158 BEECs treated with  $2 \times 10^7$  CFU/mL *E. coli*. Interestingly, S100A4 protein decreased  
159 without a significant difference after LPS treatment (Fig. 4). Additionally, S100A4  
160 protein was not detected in vaginal discharge, serum, and cell supernatant by  
161 ELISA.

#### 162 4. Discussion

163 Many S100 calcium-binding protein family members have potent antimicrobial  
164 properties and are essential components of the immune response to invading  
165 pathogens during infection and inflammation [10]. For example, S100A12 can  
166 promote vascular and valve calcification to regulate vascular inflammation  
167 in non-oxidized low-density lipoprotein-induced vascular smooth muscle cells [14].  
168 Exogenous S100A12 significantly upregulated MMP-13 and VEGF via the p38  
169 MAPK and NF- $\kappa$ B pathways [15], and high expression of S100A8/9 was observed in  
170 osteoarthritis [16].

171 While some evidences strongly support S100A4 as an inflammation-related  
172 protein, the regulatory mechanism of S100A4 is unclear [10]. One study reported that  
173 S100A4 was upregulated in LPS-treated endometritis, and a lack of S100A4 resulted  
174 in an impaired inflammatory response in mice models of acute endometritis [11].  
175 S100A4 has also been reported to induce a network of inflammatory cytokines or  
176 promote the inflammatory response in mononuclear cells [17]. However, another  
177 study found that S100A4 decreased the release of cytokines in a high fat-diet  
178 inflammatory model [18]. Therefore, the inflammatory regulation of S100A4 may be  
179 different under various conditions, suggesting that the relative mechanisms still need  
180 further study.

181 Bovine endometritis often occurs postpartum within 60 days. However, the  
182 alteration of S100A4 in bovine endometritis is unknown. In this study, we chose cows  
183 40to 60 days postpartum that experienced the prolonged uterine inflammation [12].  
184 We first found that S100A4 mRNA expression levels and protein contents decreased  
185 in bovine endometrium with endometritis (Fig. 2), but increased in some reports [9,  
186 11]. Furthermore, combined with pathological grade, S100A4 decrease was closely  
187 related to the intensity of inflammation in bovine endometrium (Fig. 1 and Fig.2).  
188 Thus, decreased S100A4 may contribute to deterioration of endometritis in dairy  
189 cows.

190 Because endometritis is an inflammatory response of endometrial epithelial cells,  
191 an inflammatory cell model should be used to research S100A4. We successfully  
192 developed a cell model using *E. coli* or LPS stimulation and evaluated it by cytokine

193 detection (Fig. 3). In vitro results showed that cytokines increased with the dosage of  
194 bacteria or LPS, however, S100A4 was negatively correlated with the dosage of *E.*  
195 *coli* or LPS in BEECs (Fig. 4). Thus, the decrease of S100A4 was also closely related  
196 to the intensity of cellular inflammation in vitro. Although these results were not in  
197 agreement with other studies [17], we deem that bovine S100A4 may be a particular  
198 factor that can be downregulated during bovine endometritis. Additionally, S100A4  
199 protein could not be detected in vaginal discharge samples and cell supernatant by  
200 ELISA. This may result from bovine S100A4 being an intracellular protein located at  
201 the perinuclear region of the cytoplasm (<https://www.uniprot.org/uniprot/P35466>),  
202 which is different from other S100A members that are secreted as extracellular  
203 proteins. Meanwhile, it has to point out that we used a commercial human ELISA kit  
204 to detect bovine S100A4 content because both proteins have 96.04% similarity in  
205 amino acid sequence (P26447 and P35466 in the Uniprot Database). Although the  
206 result is deduced, to a large extent, bovine S100A4 alteration is closely related to  
207 endometritis progression.

208 In conclusion, the decrease of intracellular S100A4 was closely related to  
209 endometritis in dairy cows, and S100A4 is a pathogenesis-related protein of  
210 endometritis. We tentatively proposed that decreased expression of S100A4 may play  
211 an important role in the development of bovine endometritis. However, the function  
212 and mechanism of S100A4 in bovine endometritis still need further investigation.

213 **Ethics statement**

214 The present study was approved by the Institutional Animal Care and Use  
215 Committee of Lanzhou Institute of Husbandry and Pharmaceutical Sciences of  
216 Chinese Academy of Agricultural Sciences (SYXK-2014-0002).

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281 **Table 1** Primer pairs used for q-PCR

Gene name	ID	Sequence
$\beta$ -actin	NM_173979.3	F- CTCTTCCAGCCTTCCTTCCT R-GGGCAGTGATCTCTTTCTGC
S100A4	NM_174595.2	F- AAAGTGAAGGCTCCTCAGGTGT R- GTTCAGTATGGTGCTCACGTCT
TNF- $\alpha$	NM_173966.3	F-TCCAGAAGTTGCTTGTGCCT R-CAGAGGGCTGTTGATGGAGG ·
IL-1 $\beta$	NM_174093.1	F-CCTCGGTTCCATGGGAGATG R- AGGCACTGTTCCCTCAGCTTC · ·
IL-6	NM_173923.2	F-GCTGAATCTTCCAAAATGGAGG R-GCTTCAGGATCTGGATCAGTG ·
IL-10	NM_174088.1	F-CCTTGTCGGAAATGATCCAGTTTT R-TCAGGCCCGTGGTTCTCA · ·

## Figure legends

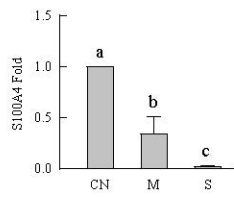
**Fig. 1** Endometrial biopsy tissue sections from dairy cows stained with HE: CN, M, and S represents endometrial biopsy from healthy cows, cows with mild endometritis, and cows with severe endometritis, respectively. Cells labeled as E, epithelial cell; L, lymphocyte; P, plasmocyte; R, red blood cell. NE represents epithelial cells necrosis. Images were magnified 400×.

**Fig. 2** Gene expression and protein content of S100A4 in endometrial tissue. Different letters mean significant difference,  $P < 0.05$ . CN, M and S represented healthy endometrium, mild endometritis and severe endometritis endometrium, respectively.

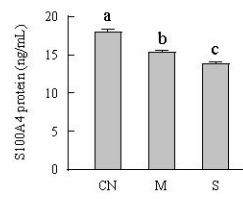
**Fig.3** Gene expression of inflammatory factors in *E. coli* or LPS-stimulated BEECs. A-D: the expression of TNF- $\alpha$ , IL-1 $\beta$ , IL-6 and IL-10 in *E. coli* stimulated BEECs. E~H: the expression of TNF- $\alpha$ , IL-1 $\beta$ , IL-6 and IL-10 in LPS stimulated BEECs. Different letters mean significant difference,  $P < 0.05$ .

**Fig. 4** Gene expression and protein content of S100A4 in *E. coli* or LPS-stimulated BEECs. Different letters mean significant difference,  $P < 0.05$ . A-B: the expression and protein contents decreased in *E. coli*-stimulated BEECs; C-D: the expression and protein contents decreased in LPS-stimulated BEECs.

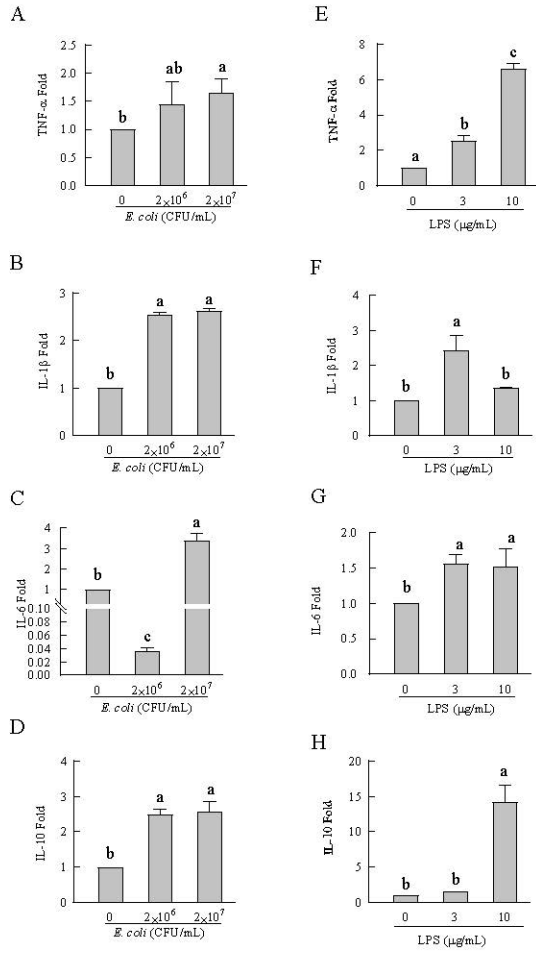
A

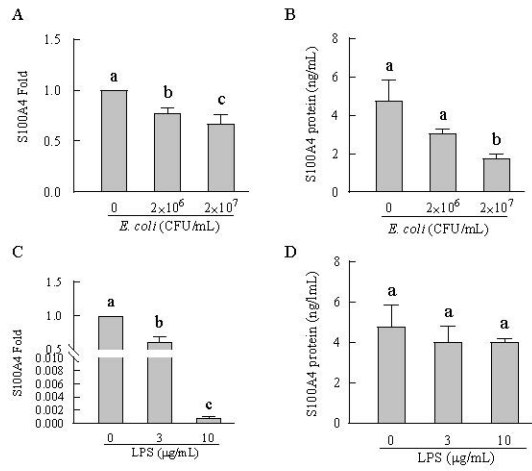


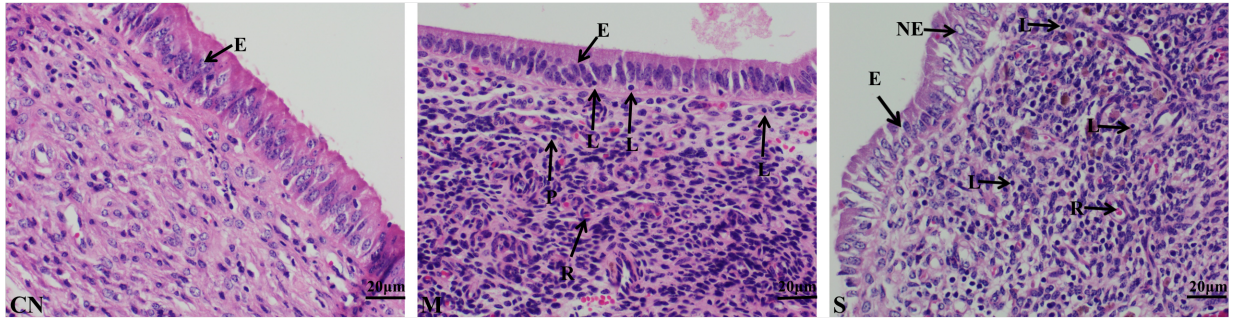
B



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1, This work firstly indicated that S100A4 decreased in endometritis of dairy cows in vivo and in vitro.

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