APPLIED MICROBIAL AND CELL PHYSIOLOGY



Sef fimbria operon construction, expression, and function for direct rapid detection of *Salmonella* Enteritidis

Xuanqiang Gu^{1,2,3} • Qianxi Hou^{1,2,3} • Jiaqi Liu^{1,2,3} • Pengpeng Xia^{1,2,3} • Qiangde Duan^{1,2,3} • Guoqiang Zhu^{1,2,3} (b)

Received: 31 March 2021 / Revised: 27 May 2021 / Accepted: 29 May 2021 / Published online: 22 June 2021 (© The Author(s), under exclusive licence to Springer-Verlag GmbH Germany, part of Springer Nature 2021

Abstract

Salmonella Enteritidis (SE) causes both horizontal and vertical transmission of diseases in poultry industry and is also one of the main causes of human food poisoning. Sequence analysis of the sef operon of poultry-derived Salmonella serotypes showed the presence of an entire sef operon in SE, whereas only sef pseudogenes were found in Salmonella Gallinarum and Salmonella Pullorum. Subsequently, the sef operon of SE was cloned into the pBR322 plasmid and expressed in a modified Escherichia coli strain SE5000. sef operon expression was demonstrated using sodium dodecyl sulfate-polyacrylamide gel electrophoresis, western blot, agglutination assay, and transmission electron microscopy. The results showed that SE5000+Sef, but not SE5000+pBR322, could specifically react with SE-positive chicken serum in an agglutination assay, which could be clearly visualized by the naked eye within less than 2 min. In contrast, SE5000+Sef could not be recognized in Salmonella Gallinarumand Salmonella Pullorum-positive chicken sera. Next, taking advantage of the exclusive presence of an entire sef operon in SE, we set up an agglutination-based detection system to monitor the dynamics of Sef-targeted antibody from SE-infected chicks for 47 days. Using the proposed detection method, SE was readily detectable starting from 2 weeks post-infection. Finally, we compared the proposed SE5000+Sef-based detection system with commercially available agglutination antigen using the classical bacterial isolation and identification procedure as reference. The results showed that the SE5000+Sef system was more consistent with the results of bacterial isolation and identification with almost 100% accuracy. We established a simple, sensitive, and cheap agglutination method for rapid and specific detection of SE-infected chickens, which can facilitate epidemiological investigation and eradication of SE infections.

Key points

• Only the Salmonella Enteritidis serotype expressed Sef fimbriae in chicken infected with SE.

• A rapid, large-scale method of detection by the naked eye of detection of SE-infected chicken is presented.

Keywords Salmonella Enteritidis · sef operon · Sef fimbria · Rapid detection

Xuanqiang Gu and Qianxi Hou contributed equally to this work and shared the first authorship.

Qiangde Duan dqd@yzu.edu.cn

Guoqiang Zhu yzgqzhu@yzu.edu.cn

- ¹ College of Veterinary Medicine, Yangzhou University, 12 East Wenhui Road, Yangzhou 225009, China
- ² Jiangsu Co-Innovation Center for Important Animal Infectious Diseases and Zoonoses, Yangzhou 225009, China
- ³ Joint International Research Laboratory of Agriculture and Agri-Product Safety of Ministry of Education of China, Yangzhou University, Yangzhou 225009, China

Introduction

Salmonella Enteritidis (SE) is a pathogen of a broad range of host organisms; it is also the main causative agent of human food poisoning and can lead to nontyphoid gastroenteritis in humans, livestock, and poultry (Gal-Mor et al. 2014). Every year, about 93.8 million people in the world are infected with Salmonella, and around 155,000 die of the Salmonella infection (Majowicz et al. 2010). In China, gastroenteritis caused by Salmonella accounts for about 40% of bacterial food poisoning. SE is the main pathogen of Salmonella food poisoning. Livestock and poultry products are the main infection sources of SE, and eggs are the primary causes of human

Salmonella infections (Gantois et al. 2009). In case of egg contamination with SE, the entire production must be discarded. For example, in 2010, 500 million "problem eggs" were recalled because of an SE outbreak in the USA (Kuehn 2010). Moreover, SE is not only the cause of human food poisoning but also the cause of great losses in livestock and poultry breeding. Therefore, diagnostic and detection methods are urgently needed as a primary step to eradicate *Salmonella* infections, especially considering that the current methods lack sensitivity and efficiency.

At present, the many types of detection methods of Salmonella include traditional biological and biochemical identification, immunological diagnosis, and molecular biological identification (Eriksson and Aspan 2007). Conventional microbiological methods include enrichment culture, preliminary culture, isolation from pure culture, and determination with a series of biochemical reactions, serological grouping, and bacterial types (Miao et al. 2017). Molecular biology protocols also have been performed for Salmonella detection; these involve nucleic acid probe detection technology (Machado et al. 2019), polymerase chain reaction (PCR) technology (Yang et al. 2020), and gene chip (Ricke et al. 2013). Other immunological diagnostic methods of Salmonella include enzyme-linked immunosorbent assay (ELISA) (Oliveira et al. 2006), Dot-ELISA (Roychoudhury et al. 2009), immunofluorescence (Stoica et al. 2018), and immunomagnetic separation (Lynch et al. 2004). These methods are limited, however, by their time-consuming and laborious nature (Li et al. 2020). Thus, optimized methods for pathogenic Salmonella detection are urgently needed.

Since the 1930s, scientists have developed a series of serological methods for detection of SE-infected chickens; these techniques include the serum tube agglutination test (STAT) (Gast 1997), latex agglutination test (LAT) (Margot et al. 2013), and agglutination test with staining antigen for whole blood tests (Yang et al. 2019). Among them, the slide agglutination test with staining antigen has the advantages of simplicity and a time-saving but moderate level of specificity. Currently, commercial agglutination antigen targeted for somatic antigen consists of the whole bacterial cell, which can be the source of nonspecific cross-reactions with many blood or serum components (Yan et al. 2011); consequently, it can be the origin of many false-positive and false-negative reactions with poor reproducibility (Fasano et al. 2017). Therefore, this method lacks both specificity and sensitivity in detection of SE-infected chickens.

Sef fimbriae are a type of fimbriae that are encoded by a *sef*ABCD operon. They are an important virulence factor involved in the initial attachment, which is an important step in the SE infection process (Zhu et al. 2009; Yue et al. 2012). Turcotte and Woodward (1993) reported that the *sef*A gene, which encodes the major subunit of Sef fimbriae, is widely conserved within *Salmonella* serotype D; the authors

developed latex particles coated with anti-SefA monoclonal antibodies as a diagnostic system to successfully detect SE and Salmonella Dublin, but not Salmonella Gallinarum and Salmonella Pullorum, which are lacking a functional Sef. Our study found that only SE had a complete sef operon among the Salmonella serotypes isolated from infected chickens. In Salmonella Gallinarum and Salmonella Pullorum, the sef operon is only represented by pseudogenes and it is completely missing in Salmonella Typhimurium. With the exception of SE, none of the Salmonella serotypes express Sef fimbriae in vivo and therefore do not induce the production of antibodies against Sef in infected chickens. On the basis of this genetic divergence between SE and other Salmonella serogroups, this study was undertaken to describe a novel agglutination method using a recombinant Escherichia coli SE5000+Sef strain (i.e., the SE5000 expressing Sef fimbriae in the optimal condition) for the detection of SE in chickens. The proposed method combines all requested qualities, such as specificity, sensitivity, rapidity, and low cost, and can be generalized as a method of choice for future SE detection in poultry farming.

Materials and methods

Bacterial strains and plasmids

The modified E. coli SE5000 strain (Schifferli et al. 1991) obtained from Dr. Dieter Schifferli (University of Pennsylvania School of Veterinary Medicine, Philadelphia, PA, USA) was stored in our laboratory (Xia et al. 2015). The standard strain C7920, used as an SE cloning strain, was purchased from China Institute of Veterinary Drug Control (Beijing, China) (Liquan Huang 2006). Salmonella Gallinarum strains, including vaccine SG9R (purchased from China Institute of Veterinary Drug Control, Beijing, China) and isolate SG01 (isolated from our laboratory, and kept in China Institute of Veterinary Drug Control, Beijing, China), and Salmonella Pullorum strains, including reference CVCC523 (purchased from China Institute of Veterinary Drug Control, Beijing, China) and isolate S08 (isolated from our laboratory, and kept in China Institute of Veterinary Drug Control, Beijing, China), were stored in our laboratory. Plasmid pBR322 purchased from New England Biolabs (NEB; Beverly, MA, USA) was used as the expression vector. All bacterial strains were grown in Luria-Bertani (LB) broth at 37 °C.

Bioinformatics analysis

We performed multiple sequence alignments using *sef* operon sequences retrieved from the National Center for

Biotechnology Information (NCBI; Bethesda, MD, USA) for different *Salmonella* Pullorum, *Salmonella* Gallinarum, and SE strains. The *sef* operon nucleotide sequence from SE strain SE95 (CP050716) was selected as a reference sequence. We used the NCBI Nucleotide BLAST tool to look for the homologous *sef* operon in *Salmonella* Pullorum, *Salmonella* Gallinarum, and SE. The presence of the *sef* operon was verified in *Salmonella* Pullorum/Gallinarum strains, including *Salmonella* Gallinarum strains (CP019035 and AM933173) and *Salmonella* Pullorum strains (CP003786, CP003047, CP006575, LK931482, CP022963, and CP012347). The sequence data were analyzed and aligned by the ClustalW method using the MEGA 7 software (version 7.0.26) and Jalview (version 2.11.0) (Waterhouse et al. 2009).

PCR, DNA cloning, and construction of expression vector

The *sef* operon was amplified from the genome of SE standard strain C7920 (NCBI: MW594401) using the upstream primer sef-up: 5'- <u>G GAT CC</u> AAA ATg gCg TgA gTA TAT TAg CAT CCg CA -3' and the downstream primer sef-lo: 5'- <u>G</u> <u>TCG AC</u>TT ATT ATA ATT CAA TTT CTG TCG CAT AT - 3'. The underlined sequences represent *Bam*HI and *Sal*I restriction enzyme sites, respectively. PCR was performed using Phanta Max Super-Fidelity DNA Polymerase (P505-d1/d2/d3, Vazyme Nanjing, China) (Ahmed et al. 2021). The PCR product of the *sef* operon and the pBR322 vector were digested by *Bam*HI and *Sal*I (NEB, Ipswich, MA). The linear DNA fragments were purified and ligated using T4 DNA ligase (NEB), and the recombinant plasmid named pBR-*sef* was confirmed by *Bam*HI and *Sal*I restriction enzyme/ electrophoresis and DNA sequencing.

Extraction of Sef fimbriae

The pBR-*sef* plasmid was electroporated into the strain SE5000; the generated strain SE5000+Sef was cultured at 37 °C for 12 h. Fimbrial protein was prepared by the hot extraction method with some changes (Khan and Schifferli 1994); in brief, strains SG9R, SG01, CVCC523, S08, and SE5000 with pBR-*sef* cultures were centrifuged at 4 °C and 4500 rpm for 10 min, and then the sediment was washed twice with sterile phosphate-buffered saline (PBS). Then, 75 mM NaCl-0.5 mM Tris-HCl (pH 7.4) solution was added, and the mixture was incubated in a water bath at 62 °C. After 30 min, the sample was centrifuged for 20 min at 8000 rpm, and 20% saturated ammonium sulfate was added to the supernatant. After overnight precipitation, the supernatant was discarded by 14,000-rpm centrifugation for 30 min, and precipitated fimbriae were resuspended in sterile PBS.

Agglutination test

pBR322 and pBR-*sef* were electroporated into the SE5000 strain to generate SE5000+pBR322 and SE5000+Sef, respectively. Positive clones were screened on plates with added ampicillin. Strains SG9R, SG01, CVCC523, S08, SE5000+pBR322, and SE5000+Sef were grown at 37 °C; the amount of bacteria was adjusted to 5×10^9 colony-forming units (CFU)/mL and washed three times with sterile saline before subjecting the sample to agglutination with the laboratory-made anti-SefA monoclonal antibody (Zhu et al. 2010). Next, 10 µL of bacterial solution was mixed on a clean glass plate with the same volume of 1:40 diluted monoclonal antibody; after a 2-min incubation, the agglutination results could be seen by the naked eye.

Sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and western blot

We used SDS-PAGE and western blot methods (Kurien and Scofield 2015) to identify the expression of Sef. Briefly, hotextracted Sef fimbriae and recombinant SefA protein (including a $6 \times$ His tag) were run onto a SDS-PAGE, followed by electro-transfer onto a nitrocellulose membrane (Biosharp, Anhui, China) using the BIO-RAD transfer system (Shanghai, China). After overnight blocking at 4 °C with 10% bovine serum albumin (BSA), the membrane was washed three times with PBS-Tween 20 (PBST) and incubated for 1 h at 37 °C with anti-SefA monoclonal antibody (diluted 1:1000 by PBST). A second washing step with PBST was made, followed by a 2-h incubation at 37 °C with a sheep anti-mouse-HRP conjugated IgG (Novus, Gudensberg, Germany). We performed a final washing step with PBST before detection with diaminobenzidine.

Transmission electron microscopy (TEM)

SE5000+pBR322 and SE5000+Sef were cultured for 12 h, and then 1 mL of each culture was centrifuged for 10 min at 4000 rpm, washed three times with the same amount of PBS, and resuspended. These samples were observed by TEM (Yang et al. 2015); briefly, a 50- μ L sample was placed on a copper grid at room temperature for 15 min. The excess bacterial solution of the copper net was absorbed by a filter paper (Whatman-Xinhua, Hangzhou, China), and the copper mesh was floated on the surface of a 2% phosphotungstic acid (pH = 7.0) dye droplet. Then, we absorbed the excess dye solution and observed the sample under TEM.

Laboratory animal and animal welfare

The specific-pathogen-free (SPF) chickens used in this study were purchased from the Comparative Medical Center of Yangzhou University. All experiments were conducted in accordance with the local ethical guidelines and national legislation.

Sample collection

Sera of SPF chickens infected with *Salmonella* Gallinarum, *Salmonella* Pullorum, *Salmonella* Typhimurium, or SE were provided both by the China Institute of Veterinary Drug Control and by experimentally infected chickens in our laboratory. Clinically positive chicken sera of *Salmonella* Gallinarum, *Salmonella* Pullorum, *Salmonella* Typhimurium, and SE were confirmed as diagnostic serum samples from farms located in 10 different provinces of China. *E. coli* O1–, *E. coli* O2–, and *E. coli* O78–infected SPF chicken sera were obtained from Dr. Song Gao. A total of 160 sera samples were used in this study; the sources of the sera are shown in Table 1.

Specifically targeted detection for SE

To prepare SE5000+Sef agglutination antigen and SE5000+ pBR322 vector as a negative control, we used a total of 160 sera samples of experimentally and clinically infected chicken (Table 1). Of the 160 samples with results provided in less than 2 min, 30 were SE-infected SPF chicken sera, 30 were SE clinically infected chicken sera, and each of the 10 SPF chicken sera samples was infected by *Salmonella* Gallinarum, *Salmonella* Pullorum, and *Salmonella* Typhimurium; each of the 10 positive chicken sera was clinically infected by

 Table 1
 The sources of sera analyzed in this study

Salmonella Gallinarum, Salmonella Pullorum, and Salmonella Typhimurium with experimental diagnosis; and 10 were SPF chicken sera. We also included O1, O2, and O78 *E. coli* serotype–positive SPF chicken sera.

Sensitively targeted detection for SE

Ten SPF chicks at the age of 5 days were infected orally with SE strain C7920, with the infection dose of 10^7 CFU in 100 μ L of sterile LB broth; 10 SPF chicks were treated with the same volume of LB as control. Sera of SPF chicken were then collected every week and were subjected to agglutination testing by the SE5000+Sef system until positive detection of SefA-specific antibodies.

Isolation and identification of Salmonella strains

Salmonella strains were isolated from the tissue samples of chickens using the Chinese National Standard method (GB 4789.4–2010), with some optimized modifications. Briefly, each sample (1 g) was added to 10.0 mL selenite cystine broth (SC; Qingdao Hope Bio-Technology Ltd., Qingdao, China) and incubated at 37 °C; after 24 h of incubation, 5 μ L of each culture was streaked onto xylose lysine deoxycholate selective medium (XLD; Qingdao Hope Bio-Technology Ltd.) plates and incubated at 37° C for 24 h. Next, suspected Salmonella colonies were further identified by PCR assays using universal primers for *flgE* (the upstream primer of *flgE*: ACGGACCC TGTACCGTCTAAA; the downstream primer of *flgE*: TGATGTTCACCGTACCGCC) for detecting all

	Source	<i>Salmonella</i> Enteritidis Number of sera	<i>Salmonella</i> Gallinarum Number of sera	<i>Salmonella</i> Pullorum Number of sera	<i>Salmonella</i> Typhimurium Number of sera
Experimental infection	This lab	10	0	0	0
	China Institute of Veterinary Drug Control	20	10	10	10
Diagnostic serum	Jiangsu	15	2	0	0
samples	Zhejiang	10	1	0	0
	Anhui	0	5	0	0
	Henan	5	0	0	0
	Shandong	0	1	0	0
	Hebei	0	1	0	0
	Shanxi	0	0	0	1
	Hunan	0	0	5	0
	Guangxi	0	0	5	0
	Jilin	0	0	0	9
Total		60	20	20	20

Note: 10 of the SPF chicken sera produced by our laboratory and 10 of each *E. coli* O1–, *E. coli* O2–, and *E. coli* O78–infected SPF chicken sera obtained from Dr. Song Gao were not listed here

Salmonella species (Yang et al. 2020), which can conveniently and rapidly detect flagellated and nonflagellated Salmonella spp. sdf1 primers (the upstream primer of sdf1: TGTGTTTTATCTGATGCAAGAGG; the downstream primer of sdf1: TGAACTACGTTCGTTCTTCTGG) were used to identify the SE serotype (Kasturi and Drgon 2017), and glgC primers (the upstream primer of glgC: CGGTGTAC TGCCCGCTAT; the downstream primer of glgC: CTGGGCATTGACGCAAA) were used to identify the Salmonella Pullorum serotype (Kang et al. 2011).

Data analysis

In this study, we tested the sensitivity and specificity of the calculations using an online calculator (https://www. medcalc.org/calc/diagnostic_test.php). The number of true positives (a), false negatives (b), false positives (c), and true negatives (d) was entered into the four fields of the website, and the values of sensitivity, specificity, and corresponding 95% confidence interval were obtained.

Results

Pseudogenes found of the *sef* operon of *Salmonella* Gallinarum and *Salmonella* Pullorum in infected poultry

We used the NCBI database to compare the sef operon of SE with two strains of Salmonella Gallinarum and six strains of Salmonella Pullorum. The results showed the presence of substitutions and insertions in Salmonella Gallinarum CP019035 and AM933173; these mutations corresponded to either stop codons or frameshifts within the sef operon. The substitution of G to T at nt 1626 (counted from the transcriptional start site) of the sef operon caused a replacement of a glutamic acid codon with a stop codon, resulting in a truncated sefC. Other nucleotide substitutions, insertions, and deletions were found in Salmonella Pullorum CP003786, CP003047, CP006575, LK931482, CP022963, and CP012347. In the CP003786, CP003047, CP006575, and LK931482 strains, a mutation of G to A at nt 2512 (counted from the transcription start site) caused a tryptophan codon to stop codon substitution, resulting in early termination of sefC. We also noticed the presence of an insertion within the sefC open-reading frame in CP012347, in addition to the absence of a part of sefC and the entire sefD sequences in the CP022963 chromosome. Thus, we concluded that the sef operon was mainly represented by pseudogenes in Salmonella Gallinarum and Salmonella Pullorum (Fig. 1).

Amplification, cloning, and identification of the *sef* operon in SE

We used *sef*-specific primers to amplify the entire *sef* operon. The generated PCR product migrated as a DNA band of about 4268 bp in a 1% agarose gel (Fig. 2A), which was in agreement with the predicted size. The *sef* PCR product was cloned into the pBR322 plasmid generating a recombinant vector of 8212 bp (Fig. 2B).

Verification of the expression of Sef fimbriae

Upon transformation of the SE5000 strain with recombinant pBR322 harboring sef, we confirmed the expression of Sef fimbriae in the generated strain using an agglutination assay. The results showed a significant agglutination with the laboratory-made anti-SefA monoclonal antibody. In contrast, the SE5000 strain transformed with the empty vector did not react with the monoclonal antibody during the agglutination test (Fig. 3). We noticed that the SG9R, SG01, CVCC523, and S08 strains could not recognize the anti-SefA monoclonal antibody (results not shown). To confirm the expression of Sef, fimbriae were extracted from the SE5000 recombinant strain and subjected to both SDS-PAGE and immunoblotting using anti-SefA-specific antibody. As shown in Fig. 4, a visible band of about 14.3 kDa was detected in both Coomassie staining and western blot gels (Fig. 4A), which was consistent with the predicted size of SefA, the main subunit of fimbriae. There were no bands at 14.3 kDa by both SDS-PAGE and immunoblotting for the fimbriae extracted from the SG9R, SG01, CVCC523, and S08 strains (results not shown). For further confirmation, we subjected the SE5000 strains carrying either a pBR322-sef or the empty pBR322 to TEM analysis; we found obvious fimbria structures on the surface of strain SE5000+Sef but not on the negative control SE5000+ pBR322 (Fig. 5).

SE5000+Sef system used to specifically detect SEinfected chickens

We took advantage of the genetic difference between SE and other *Salmonella* serogroups to design a diagnosis system consisting of the generated strains SE5000+pBR322 and SE5000+Sef as the reporter strains for agglutination assays to detect antifimbriae antibodies in serum prepared from infected chicken. We conducted a proof-of-concept study in which we tested our designed system against 160 serum samples from experimentally and clinically infected chicken. The results showed that the negative control SE5000+pBR322 did not react with any serum sample, whereas SE5000+Sef exhibited significant agglutination with 30 SPF chicken sera infected with SE and with 30 clinical SE-infected sera, with a sensitivity of 100% (95% confidence interval: 94.04% to 100%)



Fig. 1 Mutations of the sef operon in Salmonella Gallinarum and Salmonella Pullorum genomes. Red rectangles indicate the locations of the mutations

and specificity of 100% (95% confidence interval: 96.38% to 100%). Moreover, we did not observe a cross-reaction between SE5000+Sef and samples prepared from chicken infected with *Salmonella* Gallinarum, *Salmonella* Pullorum, *Salmonella* Typhimurium, and *E. coli* (Table 2). Taken together, these data demonstrated the powerful specificity of our diagnosis tool using the SE5000+Sef system.

Infected chickens can be detected as early as 2 weeks after infection by SE5000+Sef in the agglutination assay system

Weekly antibody surveillance of SPF chicks infected with SE showed that the antibody dynamics were in accordance with the general antibody production by antigen induction (Fig. 6).



Fig. 2 (**A**) PCR amplification results of the *sef* operon. Lane M, Trans 2K plus II DNA marker (TransGen Biotech, Beijing, China); lane 1, PCR amplification product of the *sef* operon; lane 2, SE5000 control. (**B**) The result of restriction digestion of the pBR-*sef* plasmid. Lane M, Trans 15K

DNA marker (TransGen Biotech, Beijing, China); lane 1, circular pBRsef plasmid; lane 2, pBR-sef linearized by digestion with SalI; lane 3, pBR-sef linearized by digestion with XbaI



Fig. 3 The agglutination reaction results of SE5000+pBR322 and SE5000+Sef with the monoclonal antibody against SefA. (A) SE5000+pBR322. (B) SE5000+Sef

In other words, we did not detect any antibody against Sef in the first week (7 days after infection). However, 2 weeks postinfection, we detected anti-Sef antibodies in two of the 10 individuals. At 6 weeks post-infection, all of the infected SPF chicks were detected positive compared with the noninfected SPF chicks.

The SE5000+Sef system has better sensitivity and specificity than commercial agglutination antigen

For further validation of the robustness of our designed diagnosis system, we performed a comparison between a commercially available agglutination antigen and the SE5000+Sef system. The assay included subjecting forty 60-day-old laying hens selected from a chicken farm in Guangxi province to both the commercial *Salmonella* Pullorum–based and the SE5000+Sef system agglutination tests, in parallel to the standard bacterial isolation and identification method. Using the commercial agglutination kit, half of the population was positive and the other half was negative. Only nine SE and three *Salmonella* Pullorum infections of the 40 samples were

Fig. 4 (**A**) SDS-PAGE and (**B**) western blot analysis of Sef fimbriae. Lane M, molecular weight protein marker (Baiaolab, Beijing, China); lane 1, extracted Sef fimbria subunit; lane 2, recombinant SefA subunit as a control





Fig. 5 Transmission electron microscopy images. (**A**) The strain SE5000+pBR322 as a negative control only containing the pBR322 vector; (**B**) the strain SE5000+Sef expressing Sef fimbriae (black arrows point to fimbriae)

isolated from mixed tissue samples of chickens using a classical bacterial isolation procedure. Data obtained from using the SE5000+Sef system showed that 10 positive samples were detected as SE-infected chickens and nine of them were consistent with the result of the bacterial isolation and identification method. Unlike the data from SE5000+Sef, however, the results of the commercial Salmonella Pullorum-based agglutination antigen did not match with the results of bacterial isolation and identification (Table 3). Taking the bacterial isolation and identification method as a standard, the sensitivity of the SE5000+Sef system was 100% (95% confidence interval: 66.37% to 100%) and the specificity was 96.88% (95%) confidence interval: 83.78% to 99.92%). Thus, the SE5000+ Sef system had a much better sensitivity and specificity than the commercial agglutination antigens that are still widely utilized as the main detection method.

Discussion

Sef fimbriae were believed to be unique to *Salmonella* serotype D. Thorns et al. (1990) first detected the expression of Sef fimbriae in SE and *Salmonella* Dublin isolates; then, they reported that Sef was expressed by organisms only within group D serotypes of SE, including *Salmonella* Dublin and very rare *Salmonella* Blegdam and *Salmonella* Moscow serovars. A similar conclusion was reported by Turcotte and Woodward (1993), who found that the *sefA* gene was also



Samples	Agglutination with SE5000+ pBR322 (positive / total)	Agglutination with SE5000+Sef (positive / total)
SPF chicken sera	0/10	0/10
SPF chicken sera infected by Salmonella Enteritidis	0/30	30/30
SPF chicken sera infected by Salmonella Gallinarum	0/10	0/10
SPF chicken sera infected by Salmonella Pullorum	0/10	0/10
SPF chicken sera infected by Salmonella Typhimurium	0/10	0/10
Clinical Salmonella Enteritidis-positive chicken sera	0/30	30/30
Clinical Salmonella Gallinarum-positive chicken sera	0/10	0/10
Clinical Salmonella Pullorum-positive chicken sera	0/10	0/10
Clinical Salmonella Typhimurium-positive chicken sera	0/10	0/10
Clinical E. coli O1-positive chicken sera	0/10	0/10
Clinical E. coli O2-positive chicken sera	0/10	0/10
Clinical E. coli O78-positive chicken sera	0/10	0/10
Total	0/160	60/160

Table 2 Application of the SE5000+Sef detection system in analyses of 160 different chicken sera

conserved in Salmonella Gallinarum and Salmonella Pullorum, but it was not clear why Sef was not expressed in Salmonella Gallinarum and Salmonella Pullorum. According to bioinformatics analysis, the sefC and sefD genes of the sef operon in Salmonella Gallinarum and Salmonella Pullorum variably harbor different point mutations, a fragment insertion, and a deletion. The SE5000+Sef detection system did not agglutinate with antibodies produced in chickens infected both naturally and experimentally by Salmonella Gallinarum and Salmonella Pullorum, and no Sef fimbriae could be detected from the extracts of Salmonella Gallinarum and Salmonella Pullorum strains, which was consistent with our



Fig. 6 The result of the SE5000+Sef system monitoring *Salmonella* Enteritidis–infected SPF chickens. Ten 5-day-old SPF chicks were infected with *Salmonella* Enteritidis (the infection group), while the control group included 10 SPF chicks matched for age and treated by LB as control. Antibodies of the infected chicks were tested every week, infected chickens could be detected after 2 weeks, and all infected chicks could be detected until the 6th week after infection

bioinformatics data analysis showing the presence of pseudogenes in the *sef* operon of these strains.

At present, the main serotypes of Salmonella isolates from chicken flocks are Salmonella Gallinarum and Salmonella Pullorum, SE, and Salmonella Typhimurium (Wang et al. 2020). Unlike both Salmonella Gallinarum and Salmonella Pullorum, which are isolated from sick chickens, SE exists only in healthy-looking chicken (Wang et al. 2020). Importantly, the traditional serotyping is determined by somatic antigen O, and SE cannot be differentiated from Salmonella Gallinarum and Salmonella Pullorum because they share the same O antigens (O1, O9, and O12) (Yang et al. 2019). Therefore, specific and rapid detection methods for SE in chicken are very important. At present, although laboratory-manipulated methods include microbial isolation and identification, PCR and other molecular biology methods, and ELISA and other immunological methods, they all fail to perform as a rapid and low-cost screening for SE in large-scale chicken farms, or in eradicating SE (Eriksson and Aspan 2007). The traditional agglutination method has the advantages of being fast and economical, and it can be used for large-scale detection of pathogens on the spot; however, it has been criticized for being less specific and less sensitive because of confusing false-positive and false-negative results (Schreier et al. 2013). This confusion is due to the fact that commercial agglutination antigens use somatic antigens that are produced by all Salmonella species. Therefore, the diversity and complexity of the used antigens are a source of undesired nonspecific cross-reactions with different chicken sera, especially with obvious cross-reactions clinically by certain Enterobacteriae-infected chicken sera (Yang et al. 2019). We also noticed that the traditional agglutination method detected somatic antigen-based antibodies with low sensitivity because of the weak immunogenicity of the somatic antigen.

Table 3	Comparison of the res	ults of three differen	t detection tests fo	r Salmonella	Enteritidis applie	d to 40 different chicken
---------	-----------------------	------------------------	----------------------	--------------	--------------------	---------------------------

Samples	Isolation and identification of bacteria	Commercial agglutination antigen	SE5000+pBR322	SE5000+Sef
N1	/	-	_	-
N2	/	-	-	-
N3	/	-	-	-
N4	/	-	-	-
N5	/	-	-	-
N6	/	-	-	-
N7	SE	-	-	+
N8	/	-	-	-
N9	/	-	-	-
N10	/	-	-	-
N11	/	-	-	-
N12	/	-	-	-
N13	/	-	-	-
N14	/	-	-	-
N15	/	-	-	-
N16	/	-	-	-
N17	/	-	-	-
N18	/	-	-	-
N19	/	-	-	-
N20	/	_	-	-
P1	/	+	-	+
P2	SE	+	-	+
P3	/	+	-	-
P4	SE	+	-	+
P5	/	+	-	-
P6	SE	+	-	+
P7	/	+	-	-
P8	SE	+	_	+
P9	SE	+	_	+
P10	SE	+	-	-
P11	/	+	_	-
P12	SP	+	-	-
P13	/	+	-	+
P14	SP	+	_	-
P15	/	+	_	-
P16	SE	+	_	+
P17	SE	· +	-	+
P18	/	+	-	-
P10	, SP	_	_	-
P20	/	+	-	-
Total	9/40	20/40	- 0/40	-
1 Otal	2/TU	20/TU	0/40	10/40

"SE," isolated and identified Salmonella Enteritidis from chicken; "SP," isolated and identified Salmonella Pullorum from chicken; "/," not isolated; "+," positive agglutination reaction; "-," negative agglutination reaction

Therefore, an alternative method for chicken SE detection is urgently needed to meet the requirements of a rapid, specific, sensitive, and economic agglutination detection test.

Considering the unique expression of Sef fimbriae in SE among *Salmonella* serotype D, Sef can be used for exclusive

detection of SE in infected poultry by serological methods. For instance, a monoclonal antibody–coated latex reagent against the major subunit SefA (molecular weight of 14.3 kDa) was used to identify SE but came with a high number of other bacteria (Thorns et al. 1994). Rajashekara et al. (1998)

improved LAT by using recombinant SefA protein instead of monoclonal antibody-coated latex particles for detecting anti-SefA antibodies. Their method was not tested against SEinfected samples, however, and little is known about the sensitivity of the developed LAT method. We wondered whether an alternative method could be designed to detect patients or sick animals infected with SE. In this study, we cloned the sef operon of SE and confirmed its expression in SE5000. We then validated the proof-of-principle of the designed SE5000+Sef system for SE detection. Unlike the classical agglutination and LAT methods, our SE5000+Sef detection system included an internal negative control, SE5000+ pBR322, with zero detectable agglutination with serum samples tested. We next tested the efficiency of the SE5000+Sef detection system; the obtained results showed that the infected SPF chicks could be detected relatively early by the SE5000+ Sef system, starting from 2 weeks post-infection. In addition, the detection system showed strong specificity and sensitivity compared with the commercial Salmonella Pullorum-based agglutination antigens, and the results were completely consistent with the results of bacterial isolation and identification.

By optimizing the traditional plate agglutination reaction with much more specificity and sensitivity, this study proposed a method for large-scale, rapid, and on-site detection of SE in chickens. Considering its convenience for epidemiological investigation and disease control and prevention of SE in chickens, our Sef fimbriae–based method is suggested as a promising and robust technique that can be generalized for future SE detection in poultry farming.

Acknowledgements The authors gratefully acknowledge Professor Song Gao in Yangzhou University, College of Veterinary Medicine, for providing the serum for this study.

Author contribution XG and QH performed the experiments, analyzed the data, and wrote the manuscript. JL and PX participated in the data analysis and wrote the paper. QD and GZ conceived and designed the study, participated in experimental work, and wrote the paper. All authors read and approved the final manuscript.

Funding This work was supported by the College Students Research & Practice Innovation Program, Graduates Research & Practice Innovation Program of Jiangsu Province (No. KYCX20_3004), International Collaboration Program from Science and Technology Agency of Jiangsu Province (2019), and Priority Academic Program Development of Jiangsu Higher Education Institutions.

Data availability The authors declare that the data supporting the findings of this study are available within the article.

Declarations

Ethics approval and consent to participate This article does not contain any studies with human participants performed by any of the authors. The current study was approved by the Institutional Animal Care and Use Committee of the Yangzhou University College of Veterinary Medicine of China.

Consent for publication All authors listed on this manuscript have read and agreed to the publication of this research.

Competing interests The authors declare no competing interests.

References

- Ahmed A, Wang M, Khan R, Shah AA, Guo H, Malik S, Xia K, Hu Z (2021) A splice-site variant (C.3289-1G>T) in OTOF underlies profound hearing loss in a Pakistani kindred. BMC Med Genet 14:2. https://doi.org/10.1186/s12920-020-00859-x
- Eriksson E, Aspan A (2007) Comparison of culture, ELISA and PCR techniques for *Salmonella* detection in faecal samples for cattle, pig and poultry. BMC Vet Res 3:21. https://doi.org/10.1186/1746-6148-3-21
- Fasano RM, Sullivan HC, Bray RA, Gebel HM, Meyer EK, Winkler AM, Josephson CD, Stowell SR, Duncan A, Roback JD (2017) Genotyping applications for transplantation and transfusion management: the Emory experience. Arch Pathol Lab Med 141(3): 329–340. https://doi.org/10.5858/arpa.2016-0277-SA
- Gal-Mor O, Boyle EC, Grassl GA (2014) Same species, different diseases: how and why typhoidal and non-typhoidal Salmonella enterica serovars differ. Front Microbiol 391(5):1–10. https://doi. org/10.3389/fmicb.2014.00391
- Gantois I, Ducatelle R, Pasmans F, Haesebrouck F, Gast R, Humphrey TJ, Van Immerseel F (2009) Mechanisms of egg contamination by *Salmonella* Enteritidis. FEMS Microbiol Rev 33(4):718–738. https://doi.org/10.1111/j.1574-6976.2008.00161.x
- Gast RK (1997) Detecting infections of chickens with recent *Salmonella* Pullorum isolates using standard serological methods. Poult Sci 76(1):17–23. https://doi.org/10.1093/ps/76.1.17
- Kang M-S, Kwon Y-K, Jung B-Y, Kim A, Lee K-M, An B-K, Song E-A, Kwon J-H, Chung G-S (2011) Differential identification of *Salmonella enterica* subsp *enterica* serovar Gallinarum biovars Gallinarum and Pullorum based on polymorphic regions of *glgC* and *speC* genes. Vet Microbiol 147(1-2):181–185. https://doi.org/ 10.1016/j.vetmic.2010.05.039
- Kasturi K, Drgon T (2017) Real-time PCR method for detection of Salmonella spp. in environmental samples. Appl Environ Microbiol 83(14). https://doi.org/10.1128/aem.00644-17
- Khan AS, Schifferli DM (1994) A minor 987P protein different from the structural fimbrial subunit is the adhesin. Infect Immun 62(10): 4233–4243. https://doi.org/10.1128/iai.62.10.4233-4243
- Kuehn BM (2010) Salmonella cases traced to egg producers findings trigger recall of more than 500 million eggs. JAMA-J AM Med Assoc 304(12):1316. https://doi.org/10.1001/jama.2010.1330
- Kurien BT, Scofield RH (2015) Western blotting: an introduction: methods and protocols. Methods Mol Biol 1312:17–30
- Li J, Ma B, Fang J, Zhi A, Chen E, Xu Y, Yu X, Sun C, Zhang M (2020) Recombinase polymerase amplification (RPA) combined with lateral flow immunoassay for rapid detection of *Salmonella* in food. Foods 9(27):1–12. https://doi.org/10.3390/foods9010027
- Liquan Huang YW (2006) Study on the antibacterial activity of Leontopodium leontopodioides (Willd.) Beauv. in vitro. J Traditional Chinese Vet Med 1:5–7. https://doi.org/10.13823/j. cnki.jtcvm.2006.01.001
- Lynch MJB, Leon-Velarde CG, McEwen S, Odumeru JA (2004) Evaluation of an automated immunomagnetic separation method for the rapid detection of *Salmonella* species in poultry

environmental samples. J Microbiol Methods 58(2):285–288. https://doi.org/10.1016/j.mimet.2004.04.005

- Machado I, Garrido V, Hernandez LI, Botero J, Bastida N, San-Roman B, Grillo M-J, Hernandez FJ (2019) Rapid and specific detection of *Salmonella* infections using chemically modified nucleic acid probes. Anal Chim Acta 1054:157–166. https://doi.org/10.1016/j. aca.2018.12.027
- Majowicz SE, Musto J, Scallan E, Angulo FJ, Kirk M, O'Brien SJ, Jones TF, Fazil A, Hoekstra RM, Int Collaboration Enteric Dis B (2010) The global burden of nontyphoidal *Salmonella* gastroenteritis. Clin Infect Dis 50(6):882–889. https://doi.org/10.1086/650733
- Margot H, Saegesser G, Zweifel C, Stephan R (2013) Performance of a commercially available latex agglutination test kit for the identification of *Salmonella*. J Food Saf Food Qual 64(3):75–77. https://doi. org/10.2376/0003-925x-64-75
- Miao J, Han N, Qiang Y, Zhang T, Li X, Zhang W (2017) 16SPIP: a comprehensive analysis pipeline for rapid pathogen detection in clinical samples based on 16S metagenomic sequencing. BMC Bioinformatics 18:255–259. https://doi.org/10.1186/s12859-017-1975-3
- Oliveira GH, Berchieri A Jr, Montassier HJ (2006) Chicken serologic response to *Salmonella enterica* serotype Typhimurium assessed by ELISA. Braz J Poult Sci 8(1):51–54. https://doi.org/10.1590/ s1516-635x2006000100008
- Rajashekara G, Munir S, Lamichhane CM, Back A, Kapur V, Halvorson DA, Nagaraja KV (1998) Application of recombinant fimbrial protein for the specific detection of *Salmonella* Enteritidis infection in poultry. Diagn Microbiol Infect Dis 32(3):147–157. https://doi.org/ 10.1016/s0732-8893(98)00091-1
- Ricke SC, Khatiwara A, Kwon YM (2013) Application of microarray analysis of foodborne *Salmonella* in poultry production: a review. Poult Sci 92(9):2243–2250. https://doi.org/10.3382/ps.2012-02740
- Roychoudhury P, Murugkar HV, Rahman H, Subudhi PK (2009) Dot-ELISA for detection of enterotoxin production by *Salmonella* isolates from Japanese quail. Indian J Anim Sci 79(2):161–163
- Schifferli DM, Beachey EH, Taylor RK (1991) 987P fimbrial gene identification and protein characterization by T7 RNA polymerase induced transcription and *TnphoA* mutagenesis. Mol Microbiol 5:61– 70
- Schreier S, Doungchawee G, Chadsuthi S, Triampo D, Triampo W (2013) Leptospirosis: current situation and trends of specific laboratory tests. Expert Rev Clin Immunol 9(3):263–280. https://doi.org/ 10.1586/eci.12.110
- Song Q, Shen X, Yang Y, Zhang D, Gao H (2016) Genetically similar isolates of *Salmonella enterica* serotype Enteritidis persistent in China for a long-term period. J Food Sci 81(7):M1778–M1781. https://doi.org/10.1111/1750-3841.13339
- Stoica C, Ancuta PN, Lucaciu IE, Banciu AR, Sorea S, Atanasescu A, Lazar MN (2018) Computerized high-tech detection technology of immunofluorescence labelled waterborne pathogenic bacteria. Rev Chim-Bucharest 69(11):4166–4170
- Thorns CJ, Sojka MG, Chasey D (1990) Detection of a novel fimbrial structure on the surface of *Salmonella* Enteritidis by using a monoclonal antibody. J Clin Microbiol 28(11):2409–2414. https://doi. org/10.1128/jcm.28.11.2409-2414.1990

- Thorns CJ, McLaren IM, Sojka MG (1994) The use of latex particle agglutination to specifically detect *Salmonella* Enteritidis. Int J Food Microbiol 21(1-2):47–53. https://doi.org/10.1016/0168-1605(94)90199-6
- Turcotte C, Woodward MJ (1993) Cloning, DNA nucleotide sequence and distribution of the gene encoding the Sef fimbrial antigen of *Salmonella* Enteritidis. J Gen Microbiol 139(7):1477–1485. https://doi.org/10.1099/00221287-139-7-1477
- Wang X, Wang H, Li T, Liu F, Cheng Y, Guo X, Wen G, Luo Q, Shao H, Pan Z, Zhang T (2020) Characterization of *Salmonella* spp. isolated from chickens in central China. BMC Vet Res 16(1):299–308. https://doi.org/10.1186/s12917-020-02513-1
- Waterhouse AM, Procter JB, Martin DMA, Clamp M, Barton GJ (2009) Jalview version 2-a multiple sequence alignment editor and analysis workbench. Bioinformatics 25(9):1189–1191. https://doi.org/10. 1093/bioinformatics/btp033
- Xia P, Song Y, Zou Y, Yang Y, Zhu G (2015) F4+ enterotoxigenic Escherichia coli (ETEC) adhesion mediated by the major fimbrial subunit FaeG. J Basic Microbiol 55(9):1118–1124. https://doi.org/ 10.1002/jobm.201400901
- Yan M, Tam FCH, Kan B, Lim PL (2011) Combined rapid (TUBEX) test for typhoid-paratyphoid a fever based on strong anti-O12 response: design and critical assessment of sensitivity. PLoS One 6(9):e24743. https://doi.org/10.1371/journal.pone.0024743
- Yang F-H, Zhang Q, Liang Q-Y, Wang S-Q, Zhao B-X, Wang Y-T, Cai Y, Li G-F (2015) Bioavailability enhancement of paclitaxel via a novel oral drug delivery system: paclitaxel-loaded glycyrrhizic acid micelles. Molecules 20(3):4337–4356. https://doi.org/10.3390/ molecules20034337
- Yang B, Niu Q, Yang Y, Dai P, Yuan T, Xu S, Pan X, Zhu G (2019) Selfmade Salmonella Pullorum agglutination antigen development and its potential practical application. Poult Sci 98(12):6326–6332. https://doi.org/10.3382/ps/pez453
- Yang Y, Wang P, Xia P, Yang B, Dai P, Hong T, Li J, Meng X, El Qaidi S, Zhu G (2020) Rapid detection of flagellated and non-flagellated *Salmonella* by targeting the common flagellar hook gene *flgE*. Appl Microbiol Biotechnol 104(22):9719–9732. https://doi.org/10.1007/ s00253-020-10925-0
- Yue M, Rankin SC, Blanchet RT, Nulton JD, Edwards RA, Schifferli DM (2012) Diversification of the *Salmonella* fimbriae: a model of macro- and microevolution. PLoS One 7(6):e38596. https://doi.org/ 10.1371/journal.pone.0038596
- Zhu CH, Musa HH, Wu SL, Zhu GQ (2009) The role of Sef fimbriae in pathogenesis and enhancing the immunity of *Salmonella* Enteritidis. Afr J Microbiol Res 3(5):191–194
- Zhu CH, Wu J, Chen WW, Hassan HM, Zhu GQ (2010) Difference and variation of the *sef* operon gene clusters in *Salmonella* Pullorum. J Basic Microbiol 50:S120–S123. https://doi.org/10.1002/jobm. 200900262

Publisher's note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.