

**RESEARCH ARTICLE****Detection of *Actinobacillus pleuropneumoniae* through Duplex PCR Based on ApxIA and ApxIVA Genes**

Huayuan Lin^{1,2§}, Huirong Mao^{1§}, Sufang Cheng^{1§}, Mengying Yi², Yezhao Ruan¹, Cong Wu¹, Zheng Xu¹, Guoliang Hu¹, Xiaoquan Guo¹, Huabin Cao¹, Lin Li¹, Yinhua Han³ and Ping Liu^{*1}

¹Jiangxi Provincial Key Laboratory for Animal Health, Institute of Animal Population Health, College of Animal Science and Technology, Jiangxi Agricultural University, No. 1101 Zhimin Avenue, Economic and Technological Development District, Nanchang 330045, Jiangxi, P. R. China; ²National and Regional Joint Engineering Laboratory for Medicament of Zoonosis Prevention and Control, College of Veterinary Medicine, South China Agricultural University, Guangzhou, China; ³Guangxi Center for Animal Disease Control and Prevention, Nanning 530001, Guangxi, PR China

*Corresponding author: PingLiu@163.com

ARTICLE HISTORY (18-074)

Received: February 18, 2018
Revised: April 04, 2018
Accepted: May 09, 2018
Published online: June 26, 2018

Key words:

Actinobacillus pleuropneumoniae
ApxI
ApxIV
Duplex PCR

ABSTRACT

Actinobacillus pleuropneumoniae (APP) is the bacterial pathogen of pleuropneumoniae, a severe contagious and fatal respiratory disease of swine. For the purpose of setting up an easy way to detect this bacteria in swine, a duplex polymerase chain reaction (PCR) assay was built up for the detection of APP. Two pairs of primers were designed according to the APP ApxI and ApxIV gene sequences. Using these primers, a duplex PCR method was developed to fast identify APP in different bacteria samples, such as *Enterococcus faecalis*, *Escherichia coli* from cattle and swine, *Staphylococcus aureus* and *Streptococcus*. Our results have shown that duplex PCR could be a potential molecular method for the diagnosis of APP infection in swine.

©2018 PVJ. All rights reserved

To Cite This Article: Lin H, Mao H, Cheng S, Yi M, Ruan Y, Wu C, Xu Z, Hu G, Guo X, Cao H, Li L, Han Y and Liu P, 2018. Detection of *Actinobacillus pleuropneumoniae* through duplex PCR based on apxia and apxiva genes. Pak Vet J, 38(3): 276-280. <http://dx.doi.org/10.29261/pakvetj/2018.058>

INTRODUCTION

Porcine pleuropneumoniae (PCP) is a highly fatal infectious respiratory disease of swine. Its main pathogenic bacteria are *Actinobacillus pleuropneumoniae* (APP) (To *et al.*, 2018). Acute PCP can cause serious economic losses and slow growth of infected swine (Gómez-Laguna *et al.*, 2014). It commonly occurs between age 8 and 16 weeks in pigs. Its clinical signs are characterized by increased respiratory rate, dyspnea, high fever, sneezing, coughing, and severe respiratory distress with cyanosis (Bosse *et al.*, 2002).

The Gram-negative bacterium *Actinobacillus pleuropneumoniae* is recognized as an obligate parasite in the respiratory tract of pigs (Bosse *et al.*, 2002; Antenucci *et al.*, 2017). Until now, 15 serotypes of *Actinobacillus pleuropneumoniae* have been identified which secrete four types of RTX (repeats-in-toxin) toxins (ApxIA, ApxIIA, ApxIIIA, and ApxIVA) (Shin *et al.*, 2011; Gómez-Laguna *et al.*, 2014; To *et al.*, 2016; Yee *et al.*, 2018). RTX toxins are the main factors of *Actinobacillus pleuropneumoniae*

virulence and the virulence level of each *Actinobacillus pleuropneumoniae* serotype may be directly related to the presence of Apx toxins (Park *et al.*, 2009; Hsu *et al.*, 2016; Hathroubi *et al.*, 2017). ApxIA is a strongly hemolytic and cytotoxic 105-kDa protein (Frey, 1995; Liu *et al.*, 2009; To *et al.*, 2016). It is commonly secreted by the most virulent strains of serotypes 1, 5, 9, 10 and 11 (Jansen *et al.*, 1993; Yee *et al.*, 2018). ApxIIA is also a 105-kDa moderately hemolytic and weakly cytotoxic protein secreted by all kinds of serotype strains except serotype 10 (Jansen *et al.*, 1993; Frey, 1995). ApxIIIA is a strongly cytotoxic but non-hemolytic 112.8-kDa protein secreted by the serotypes 2-4, 6, 8 and 15 with (Frey, 1995; Huang *et al.*, 2006; Liu *et al.*, 2009). ApxIVA is a 200-kDa weakly-hemolytic protein predicted from the protein sequence and secreted by all serotypes (Bosse *et al.*, 2002; Huang *et al.*, 2006).

However, due to the difference among the feeding environment, strain virulence, the immune status of animals and infecting dose, APP can cause infected pigs to develop a per-acute, acute, sub-acute or chronic form of the disease (Savoie *et al.*, 2000). The upper respiratory tract or the tonsils of the chronic form or asymptomatic

[§]These authors contributed equally to this work.

carriage pig may harbor APP which can infect other pigs in the same pigsty (Savoie *et al.*, 2000). Therefore, it is helpful to develop an effective way of the detection of virulent APP.

In this study, we intended to develop a duplex polymerase chain reaction (PCR) test for the detection and screening of the virulent APP based on the ApxIA and ApxIVA gene segments. We compared the difference in annealing temperature, concentration of primers and DNA template between the single PCR and the duplex PCR. We evaluated the specificity of this duplex PCR using strains of commonly-used clinical bacteria to ensure the validity of this assay.

MATERIALS AND METHODS

Bacterial strains, media and growth conditions: In this study, *Actinobacillus pleuropneumoniae* was purchased from China Veterinary Culture Collection Center. The other strains such as *Enterococcus faecalis* and *Escherichia coli* isolated from pig intestinal tract, *Staphylococcus aureus* and *Streptococcus* isolated from pig liver were provided by the Laboratory of Clinical Veterinary Medicine, Guangxi University. The strains of *Actinobacillus pleuropneumoniae* were cultured in 3% Tryptic soy broth (TBS) complemented with 10µg/mL of NAD. The other strains were grown in LB Broth.

DNA extraction and PCR amplification: The whole bacterial DNA for PCR amplification was extracted by DNA extraction kit. The detection primers of APXI A-F: 5'-TCGGTTCGTAGCATTAGCG-3' and APXI A-R: 5'-GACATCCCAACGCTGTTG-3' were redesigned according to the gene of *Actinobacillus pleuropneumoniae* ApxI (PCR product size: 507; GenBank accession no. AF240779.1). The detection primers of ApxIV A-F: 5'-CTTTTTTGTATAGAAGAATCA-3' and ApxIV A-R: 5'-TCGTCAATAGCGTAACAGTT-3' were designed according to the gene of *Actinobacillus pleuropneumoniae* ApxIV (PCR product size: 1092; GenBank accession no. FJ 848574.1). The single PCR reactions were set up into 20µL volume containing 10µL 2×Taq Master Mix, 1µL of each primer (2µmol/L of each primer), and 1µL of the extracted DNA template, and the 7µL of the ddH₂O. The configuration was 95°C for 3 min, then denaturation at 94°C for 30 seconds, annealing for 30 seconds, extension at 72°C for 1 min for 35 cycles, and then a final extension step 72°C for 10min in a thermocycler. Meanwhile, the optional annealing temperature, concentration of DNA and primers were carried out. Briefly, 6 temperatures at an interval of 2°C between 50°C to 60°C were used as the annealing temperatures for the ApxIA and ApxIVA amplification; 10 different concentrations of primer at an interval of 0.2ng/mL from 0.2ng/mL to 2.0ng/mL were used to select the best concentration of primers for target gene amplification; 10 different concentrations of DNA at an interval of 15µg/mL from 15µg/mL to 150µg/mL were used to examine the sensibility of the PCR reaction. The results of PCR reactions were examined by electrophoresis on 1% agarose and visualized by the GelDoc XR System (Bio-Rad, USA).

Development of a duplex PCR assay for the detection of *Actinobacillus pleuropneumoniae*: Duplex PCR

amplification conditions were 95°C for 3 min, then denaturation at 94°C for 30seconds, annealing for 30sec, extension at 72°C for 1 min for 35cycles, and then a final extension step 72°C for 10min in a thermocycler. Meanwhile, the optional annealing temperature, concentration of DNA and primer, PCR reaction system were carried out. Briefly, 6 different temperatures at an interval of 2°C from 50°C to 60°C were used as the annealing temperature; 9 different volume of PCR reaction system at an interval of 5µL from 10µL to 50µL, were used to this study; 10 different concentration of primer at an interval of 0.2ng/mL from 0.2ng/mL to were used to choose the best concentration of primer for target gene amplification; 10 different concentration of DNA at an interval of 24µg/mL from 24µg/mL to 240µg/mL were used to examine the sensibility of the ApxIA gene detection; 10 different concentration of DNA at an interval of 15µg/mL from 15µg/mL to 150µg/mL were used to examine the sensibility of the ApxIVA gene detection. The results of PCR reactions were examined by electrophoresis on 1% agarose and visualized by the GelDoc XR System (Bio-Rad, USA).

Specificity of the duplex PCR assay: The specificity of the duplex PCR was evaluated by comparing the results obtained using DNA extracts from the strains of *Enterococcus faecalis*, *Escherichia coli*, *Staphylococcus aureus* and *Streptococcus*. The duplex PCR products were purified by commercially available kit (TaKaRa, Otsu, Japan), and then sequencing analysis was conducted (Sangon Biotech, Shanghai, China).

RESULTS

Single PCR results: Different annealing temperatures had various effects on ApxIA gene amplification, whereas there was no obvious visualization difference in the ApxIVA gene products (Fig.1A and 1B). The results showed that the annealing temperature 52°C, 54°C, 56°C, 58°C were the best temperature for the ApxIA gene amplification (Fig. 1A). The results of different concentrations of primers showed that the higher the primer concentration was, the brighter the strap of PCR was (Fig. 2A and 2B). The best concentration of primer was determined to be 2.0ng/mL. The results of different concentrations of DNA showed that the higher the DNA concentration was, the brighter the strap of PCR product was in our study (Fig. 3A and B).

Duplex PCR results: The results of six different annealing temperatures showed that 58°C and 60°C are the best temperatures for the ApxIA and ApxIVA gene amplification (Fig. 4A). The results of 9 different volumes of PCR reaction systems showed that 30µL is the best for the ApxIA and ApxIVA gene amplification (Fig.4B). Under the 10 different ApxIA primer concentrations, the results showed that the concentrations between 0.6ng/mL and 2.0ng/mL are conducive to the ApxIA and ApxIVA gene amplification (Fig. 5A). In the 10 different concentrations of ApxIVA primer, the results showed that the concentration 1.0ng/mL is the best for the ApxIA and ApxIVA gene amplification (Fig. 5A). The PCR sensitivity examination showed that the DNA concentration 15µg/mL is the best concentration for the duplex PCR detection.

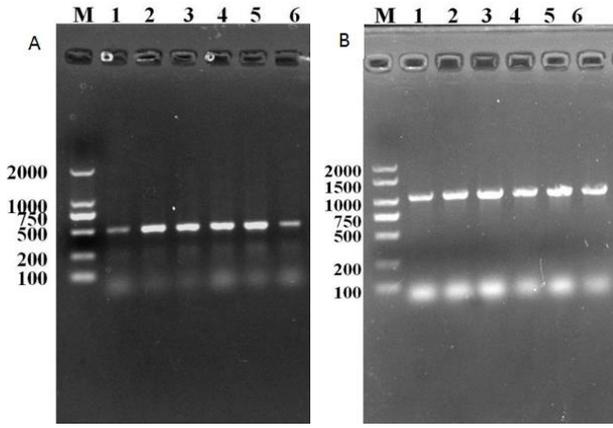


Fig. 1: Effects of 6 different annealing temperatures in ApxIA (A) and ApxIVA (B) gene amplification. M: DNA Marker 2000; 1-6: Annealing temperature 50, 52, 54, 56, 58 and 60°C.

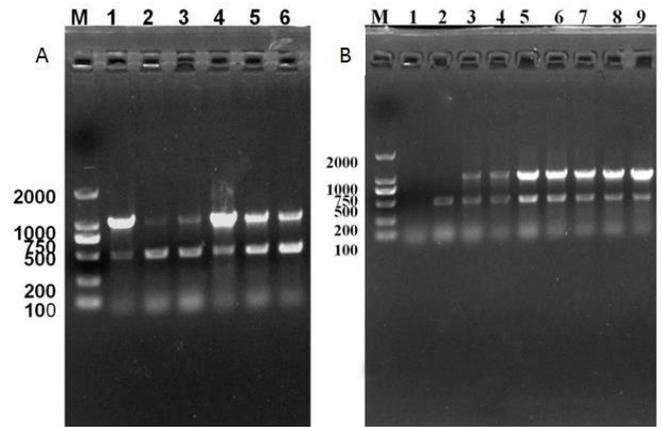


Fig. 4. A: Effects of 6 different annealing temperatures in duplex PCR amplification. M: DNA Marker 2000; 1-10: annealing temperature 50°C, 52°C, 54°C, 56°C, 58°C, 60°C. **B:** Effects of 9 different volumes of PCR reaction system in duplex PCR detection. M: DNA Marker 2000; 1-10: PCR reaction system volume 10µl, 15µl, 20µl, 25µl, 30µl, 35µl, 40µl, 45µl, 50µl. **C:** Effects of 10 different ApxI A primer concentrations in duplex PCR detection. M: DNA Marker 2000; 1-10: ApxI A primer concentration 0.2ng/mL, 0.4ng/mL, 0.6ng/mL, 0.8ng/mL, 1.0ng/mL, 1.2ng/mL, 1.4ng/mL, 1.6ng/mL, 1.8ng/mL, 2.0ng/mL.

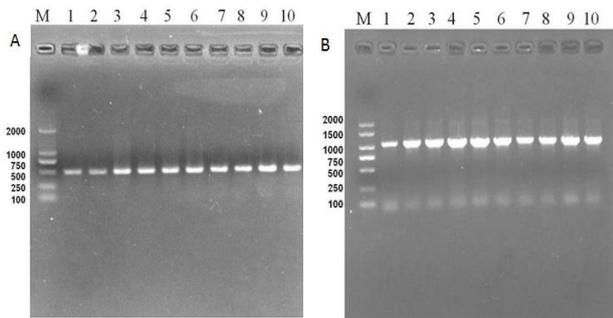


Fig. 2: Results of 10 different concentrations of ApxI A (A) ApxIV A (B) primer primer in PCR amplification. M: DNA Marker 2000; 1-10: primer concentration 0.2ng/mL, 0.4ng/mL, 0.6ng/mL, 0.8ng/mL, 1.0ng/mL, 1.2ng/mL, 1.4ng/mL, 1.6ng/mL, 1.8ng/mL, 2.0ng/mL.

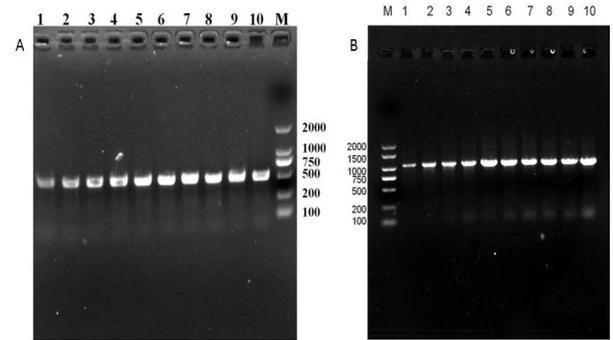


Fig. 3. A: Results of sensibility detection in ApxIA gene. M: DNA Marker 2000; 1-10: DNA concentration 24, 48, 72, 96, 120, 144, 168, 192, 216 and 240 µg/mL. **B:** Results of sensibility detection in ApxIVA gene. M: DNA Marker 2000; 1-10: DNA concentration 15, 30, 45, 60, 75, 90, 105, 120, 135 and 150 µg/mL.

Specificity of the duplex PCR assay: To test whether the duplex PCR assay can specifically amplify the ApxIA and ApxIVA gene of *Actinobacillus pleuropneumoniae*, it was used to test the strains of *Enterococcus faecalis* and *Escherichia coli*, *Staphylococcus aureus* and *Streptococcus*. The results showed that the duplex PCR assay could specifically amplify the ApxIA and ApxIVA gene of *Actinobacillus pleuropneumoniae* (Fig. 5B). The sequencing and NCBI blast analysis of the products of PCR confirmed that these sequences belong to the ApxIA and ApxIVA gene of *Actinobacillus pleuropneumoniae* (Fig. 6-7).

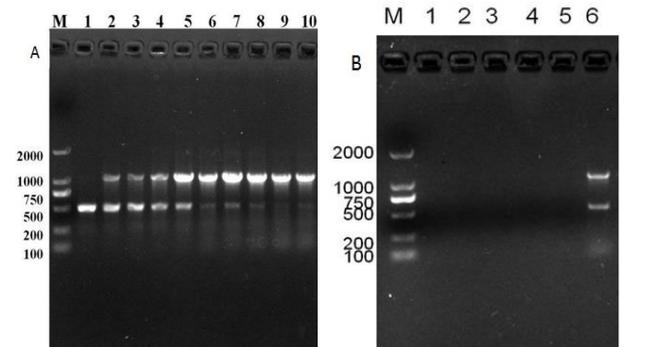


Fig. 5. A: Effects of 10 different ApxIVA primer concentrations in duplex PCR detection. M: DNA Marker 2000; 1-10: ApxIV A primer concentration 0.2ng/mL, 0.4ng/mL, 0.6ng/mL, 0.8ng/mL, 1.0ng/mL, 1.2ng/mL, 1.4ng/mL, 1.6ng/mL, 1.8ng/mL, 2.0ng/mL. **B:** Results of specificity detection in duplex PCR. M: DNA Marker 2000; 1-6: *Staphylococcus aureus*, *Escherichia coli* from cattle, *Enterococcus faecalis*, *Escherichia coli* from swine, *Streptococcus*, *Actinobacillus pleuropneumoniae*.

DISCUSSION

Previous studies have reported several techniques for the identification of *Actinobacillus pleuropneumoniae* such as ribotyping and restriction analysis of PCR(PCR-REA) (Jaglic *et al.*, 2004), multiple PCR based on the omlA gene

(Angen *et al.*, 2008) and enzyme-linked immunosorbent assay (ELISA) using the recombinant ApxIA, ApxIIA, ApxIIIA, ApxIVA proteins and lipopolysaccharide (Shin *et al.*, 2011; Sassu *et al.*, 2017; Teshima *et al.*, 2017). Tremblay *et al.* (2017) developed an ApxIV probe which could specifically detect APP in the swine lung tissue.

Apx exotoxins secreted by *Actinobacillus pleuropneumoniae* are toxic factors that deeply impact the pathogenesis of porcine pleuropneumonia. Former studies reported that ApxIVA is produced by all the 15 serotypes of APP and is highly specific to APP (Wei *et al.*, 2012; Giménez-Lirola *et al.*, 2014). Due to the specificity of the ApxIVA gene to the *Actinobacillus pleuropneumoniae* strains, the detection assays based on this gene have been widely explored for the identification of the *Actinobacillus pleuropneumoniae* strains (Schaller *et al.*, 2001; Dreyfus *et al.*, 2004; Turni and Blackall, 2007). Previous studies have reported that ApxI A may be the most virulent protein with the product in the strains could increase 100-fold of bacterial toxicity (Liu *et al.*, 2009; Hsu *et al.*, 2016; To *et al.*, 2016). Shin *et al.* (2011) studies have developed a useful ELISA for the detection of *Actinobacillus pleuropneumoniae* based on the Apx exotoxins (ApxIA, ApxII and ApxIIIA). But its major use was the detection of serum antibodies to *Actinobacillus pleuropneumoniae* and this assay was difficult to develop and apply for clinical test. Tremblay *et al.* (2017) developed a effective fluorescence in situ hybridization way using an ApxIV probe for the detection of *Actinobacillus pleuropneumoniae*. This way could specifically localize *Actinobacillus pleuropneumoniae* cells within the lung tissue by using species-specific

oligonucleotide probe. But this assay was complicated and also difficult to apply for *Actinobacillus pleuropneumoniae* clinical test. In this study, the duplex PCR assay was developed for the detection of *Actinobacillus pleuropneumoniae* strains that could simultaneously produce ApxIA and ApxIVA toxins. According to the results of single PCR (Fig.1-3), we found that the primers we designed could specifically amplify the ApxIA and ApxIVA gene segments of the *Actinobacillus pleuropneumoniae* strains. According to the results of duplex PCR, we found that an increase in ApxIVA primer concentration could induce a decrease in the ApxIA product (Fig.4B). It means that the same best reaction conditions optimized for the *single* PCR could be inappropriate to set up the *duplex* PCR. But it could help us to examine whether the primers could specifically amplify the target gene. In the duplex PCR, it is important to design the fully-optimized primers for the target gene to avoid non-specific binding among the primers. Meanwhile, the optimal template concentration, optimal annealing temperature and optimal reaction volume were also playing an important role in effectively setting up the useful duplex PCR. In this study, we also used this duplex PCR to detect other common veterinary clinical bacteria to confirm the specificity of this duplex PCR in the *Actinobacillus pleuropneumoniae* strains detection. In conclusion, the results obtained in this research show that the duplex PCR test is a specific, sensitive, and highly effective diagnostic way for simultaneous identification of *Actinobacillus pleuropneumoniae*. It could be a useful way to quickly diagnose and control the infection of *Actinobacillus pleuropneumoniae* in pigs.

	10	20	30	40	50	60	70	80	90	100	110	120
5'	TCGTCGTAGCATTAGCGATGCGTTGGATCGTTGAAGCGGCACGTAAACGTTGGTACAAATCAATGGCATTACGTTTAGCGAACGAACCTTTCGGACGCAGCGGATAACAAGGTACGGCT											120
0												
5'	GTTAAGA AACCGTGAAGACGTTCCACCGTATGGCTGAAGCTAACAAAGCGTTTGCTCACTTCCGTTGGTAAATCGTTTAGCAATATGCTTGCAGGGCTTCATCACAATAATTTGCGATGAAGCC											240
0												
5'	TTACCCTTATTTAACTCATATTAATTCCAAACAGGTAATAATATAATGGCTCGTACAACCCCTATTTACAGCTACCGTAATATTGGTATCAGTGCACATCGATGCAGGTAACA											360
0												
5'	ACTACCTCTGAGCGTATCTTATCTATACGGCGTGAGTCATAAACTTGGTGAAGTGCACGATGGTGCAGCTACTATGGACTGGATGGAACAAGAGCAAGAGCGTGGTATTACCATCACC											480
0												
5'	TCTGCTGCAACAACAGCGTGGATGTC											507
0												

Fig. 6: The sequence of ApxIA gene.

	10	20	30	40	50	60	70	80	90	100	110
5'	CTTTTTTGTATAGAAAGTACAGGCAACCGCTATATTGAAAACCTTTGGTATTGAACCTCTTGGTAAAGCAAGAAGATTTGATTTTGTGCGCGGCTTTTGGTCTAAGCTTAG										110
0											
5'	TGAATCGTGGTTTGGAAAGTATTATCGACCCATCCGGTATCGGTGGAAACCGTTAACTTTACCGCGAGGTGGAAACCTACACGTTAGACGAAACAAGGTTTAAA										220
0											
5'	GCGGAAGCGCGCAAGAAAAGCCATTGGAGTTTGTAGTGAATCGCGCGAAAGTATACGGCGGTTTAGACCAAATTTAAAAAACTATGGGACAGCGGCTCAATTAAGCATTT										330
0											
5'	ATATCAAGATAAAGATACGGGCAAAATTAACCCGATTATTTACGGCACGGCCGGCAACGACAGTAAGATTGAAGGCACTAAAATCACCCGTAGGATTGCGGGTAAAGAAG										440
0											
5'	TTACGCTTGATATTGCCAATCAGAAAATGAAAAGGCGTGTAGAGAAAATGGGGCTGCTCTGTAGTGGTTCGGATATCATTAAATGTTGTTGGAGCATTGACTCCA										550
0											
5'	ACTTTAAATAGAAATGTGCTATCACAACTTATCCAGTCTTTTTCCGATAGCTTGGCTAAACTTGATAATCCCTTAGCCCTTACACTAAAATGGCGTGGTTTATGTCAC										660
0											
5'	CGGCAAGGGGAATGATGTGCTTAAAGGAACGAACATGAGGATTTGTTTCTCGTGGTGGGGGAATGATACTTATTATCGGAGTAGGCGATACAATTAAGAGACGCCG										770
0											
5'	ACGGCAAGGTAAGTCTATTTTGTGAGAGAAAAGGGGTACCTAAGCGGATCCTAAGCGGTTAGAGTTTAGCGAGTACATAACGAAAGAAGAAATAAAGAGGTTGAA										880
0											
5'	AAGGGTTTATTAACCTACGCAGTTTTAGAAAATATAATGGGAAGAGAAAACGGCGACTTTCGCTCATGCGACTATGCTTAATGAGCTTTTTACTGATTATACTAATTA										990
0											
5'	TCGTTATGAAGTTAAAGGACTAAAATGCCCCCGTTAAAAAGTTAAAAAGTCCGTTGGTGGAGTTTACAGCTGATTATTAAGTGTACGCCCTATTGACGA										1092
0											

Fig. 7: The sequence of ApxIVA gene.

Acknowledgments: This project was supported by a grant from the National Natural Science Foundation of China awarded to PL (No. 31402266), a grant from the Natural Science Foundation of Jiangxi Province awarded to PL (No. GJJ14311) and a grant from the Technology R&D Program of Jiangxi Province awarded to PL (No. 20141BBF60035) and a grant from the Technology R&D Program of Jiangxi Province awarded to YezhaoRuan (YC2017-S185).

Authors contribution: PL, GH and XG conceived and designed the experiment. HL, HM, SC MY, HC, YH, CW, ZX, YR, LL and ZX executed the experiment of bacteria sample collection, bacteria DNA extraction, PCR amplification and gene sequence. PL and HL analyzed the data and wrote the paper; All authors interpreted the data, critically revised the manuscript for important intellectual contents and approved the final version.

REFERENCES

- Angen O, Ahrens P and Jessing SG, 2008. Development of a multiplex PCR test for identification of *Actinobacillus pleuropneumoniae* serovars 1, 7 and 12. *Vet Microbiol* 132:312-8.
- Antenucci F, Fougereux C, Bosse JT, et al., 2017. Identification and characterization of serovar-independent immunogens in *Actinobacillus pleuropneumoniae*. *Vet Res* 48:74.
- Bosse JT, Janson H, Sheehan BJ, et al., 2002. *Actinobacillus pleuropneumoniae*: Pathobiology and pathogenesis of infection. *Microbes Infect* 4:225-35.
- Dreyfus A, Schaller A, Nivollet S, et al., 2004. Use of recombinant ApxIVA in serodiagnosis of *Actinobacillus pleuropneumoniae* infections, development and prevalidation of the apxiv elisa. *Vet Microbiol* 99:227-38.
- Frey J, 1995. Virulence in *Actinobacillus pleuropneumoniae* and RTX toxins. *Trends Microbiol* 3:257-61.
- Giménez-Lirola LG, Jiang YH, Sun D, et al., 2014. Simultaneous detection of antibodies against apx toxins ApxI, ApxII, ApxIII, and ApxIV in pigs with known and unknown *Actinobacillus pleuropneumoniae* exposure using a multiplexing liquid array platform. *Clin Vaccine Immunol* 21:85-95.
- Gómez-Laguna J, Islas A, Muñoz D, et al., 2014. Infection dynamics and acute phase response of an *Actinobacillus pleuropneumoniae* field isolate of moderate virulence in pigs. *Vet Microbiol* 173:332-9.
- Hathroubi S, Loera-Muro A, Guerrero-Barrera AL, et al., 2017. *Actinobacillus pleuropneumoniae* biofilms: Role in pathogenicity and potential impact for vaccination development. *Anim Health Res Rev* 7:1-14.
- Hsu CW, Li SC and Chang NY, 2016. Involvement of NF-κB in regulation of *Actinobacillus pleuropneumoniae* exotoxin ApxI-induced proinflammatory cytokine production in porcine alveolar macrophages. *Vet Microbiol* 195:128-35.
- Huang H, Zhou R, Fan H, et al., 2006. Generation of monoclonal antibodies and epitope mapping of ApxIVA of *actinobacillus pleuropneumoniae*. *Mol Immunol* 43:2130-4.
- Jaglic Z, Svastova P, Rychlik I, et al., 2004. Differentiation of *actinobacillus pleuropneumoniae* by PCR-REA based on sequence variability of the apxIVA gene and by ribotyping. *Vet Microbiol* 103:63-9.
- Jansen R, Briaire J, Kamp EM, et al., 1993. Structural analysis of the *actinobacillus pleuropneumoniae*-RTX-toxin I (ApxI) operon. *Infect Immun* 61:3688-95.
- Liu J, Chen X, Tan C, et al., 2009. In vivo induced RTX toxin apxiva is essential for the full virulence of *Actinobacillus pleuropneumoniae*. *Vet Microbiol* 137:282-9.
- Park C, Ha Y and Kim S, 2009. Construction and characterization of an *Actinobacillus pleuropneumoniae* serotype 2 mutant lacking the Apx toxin secretion protein genes apxIIIb and apxIIIc. *J Vet Med Sci* 71:1317-23.
- Sassu EL, Bosse JT, Tobias TJ, et al., 2017. Update on *Actinobacillus pleuropneumoniae*-knowledge, gaps and challenges. *Transbound Emerg Dis* 65 S1:72-90.
- Savoie C, Jobert JL, Berthelot-Herault F, et al., 2000. A PCR assay used to study aerosol transmission of *Actinobacillus pleuropneumoniae* from samples of live pigs under experimental conditions. *Vet Microbiol* 73:337-47.
- Schaller A, Djordjevic SP, Eamens GJ, et al., 2001. Identification and detection of *Actinobacillus pleuropneumoniae* by PCR based on the gene apxIVA. *Vet Microbiol* 79:47-62.
- Shin MK, Kang ML, Cha SB, et al., 2011. An immunosorbent assay based on the recombinant ApxIa, ApxIIa, and ApxIIIa toxins of *Actinobacillus pleuropneumoniae* and its application to field sera. *J Vet Diagn Invest* 23:736-42.
- Teshima K, Lee J, To H, et al., 2017. Application of an enzyme-linked immunosorbent assay for detection of antibodies to *Actinobacillus pleuropneumoniae* serovar 15 in pig sera. *J Vet Med Sci* 79:1968-72.
- To H, Nagai S, Iwata A, et al., 2016. Genetic and antigenic characteristics of ApxIIa and ApxIIIa from *Actinobacillus pleuropneumoniae* serovars 2, 3, 4, 6, 8 and 15. *Microbiol Immunol* 60:447-58.
- To H, Teshima K, Nagai S, et al., 2018. Characterization of *Actinobacillus pleuropneumoniae* field strains antigenically related to the 3-6-8-15 group from diseased pigs in Japan and Argentina. *Rev Argentina Microbiol* 50:12-22.
- Tremblay YDN, Labrie J, Chenier S, et al., 2017. *Actinobacillus pleuropneumoniae* grows as aggregates in the lung of pigs: Is it time to refine our in vitro biofilm assays? *Microb Biotechnol* 10:756-60.
- Turni C and Blackall PJ, 2007. An evaluation of the ApxIVA based PCR-REA method for differentiation of *Actinobacillus pleuropneumoniae*. *Vet Microbiol* 121:163-9.
- Wei B, Li F, Yang H, et al., 2012. Magnetic beads-based enzymatic spectrofluorometric assay for rapid and sensitive detection of antibody against ApxIVA of *Actinobacillus pleuropneumoniae*. *Biosens Bioelectron* 35:390-3.
- Yee S, Blackall PJ and Turni C, 2018. Genetic diversity and toxin gene distribution among serovars of *Actinobacillus pleuropneumoniae* from Australian pigs. *Aust Vet J* 96:17-23.