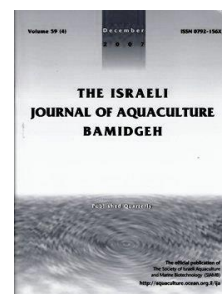




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Aeromonas hydrophila* induces apoptosis in pufferfish, *Takifugu obscurus

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Keywords: *Takifugu obscurus*; *Aeromonas hydrophila*; apoptosis; caspase-3

Abstract

Aeromonas hydrophila is a Gram-negative opportunistic pathogen causing motile aeromonad septicemia, which results in great economic losses in freshwater fish farming. In this study, we investigated the effect of *A. hydrophila* on apoptosis of pufferfish blood cells. Total blood cell count, reactive oxygen species (ROS) production, cytoplasmic free-Ca²⁺ (cf-Ca²⁺) concentration, nitric oxide (NO) production, apoptotic cell ratio and the transcription levels of caspase-3 in the pufferfish blood cells were determined after *A. hydrophila* infection. Results showed that *A. hydrophila* infection significantly depressed the total blood cell count, and significantly increased ROS production, cf-Ca²⁺ concentration, NO production and apoptotic cell ratio. The transcription levels of caspase-3 increased significantly after *A. hydrophila* infection. Taken together, our results suggest that *A. hydrophila* infection alters ROS, cf-Ca²⁺, NO levels, and active caspase-3, which lead to cell apoptosis.

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Introduction

Aeromonas hydrophila is a Gram-negative opportunistic pathogen widely distributed in aquatic environments (Majumdar et al., 2009). This bacterium is a part of normal intestinal bacterial flora of the fish and is able to cause disease through the effect of stress (Ardó et al., 2010). It causes motile aeromonad septicemia (MAS), which results in severe economic losses to the aquaculture industry worldwide (Sahu et al., 2007; Xu et al., 2012). The symptoms of *A. hydrophila* infection include anorexia, exophthalmos, dropsy, red sores, necrosis, and ulceration (Shen et al., 2013). Moreover, this bacterium can also behave as secondary opportunistic pathogens by assailing already compromised or stressed hosts (Tellez-Bañuelos et al., 2010). It can produce several virulence factors, including hemolysins, protease, leucocidins, endotoxins, and adhesions, which are all thought to contribute to the overall development of the disease in fish (Majumdar et al., 2009; Banerjee et al., 2012).

Apoptosis, also called programmed cell death, is a common physiological process that evolved in eukaryotes to remove surplus, infected, or damaged cells to maintain the integrity of the organism (Du et al., 2013). It plays a critical role in the development, cellular homeostasis, and immune responses of multicellular organisms. However, either uncontrolled apoptosis or its non-occurrence is involved in the pathogenesis of a variety of important disease processes including cancer, HIV infection, acute inflammatory disease, and autoimmunity (Thompson, 1995; Shao et al., 2004). Apoptosis also plays an important role in various bacterial diseases (Weinrauch and Zychlinsky, 1999). Induction of apoptosis in host cells appears to be part of strategies by which pathogenic bacteria modulate the immune response of the host (Moine and Abraham, 2004; Majumdar et al., 2009). *A. hydrophila* induces apoptosis in fish (Shao et al., 2004; Banerjee et al., 2012). However, the relationship between cell apoptosis and pathogenesis of fish bacterial diseases is still unclear.

Like other animals, fish blood also plays an essential role in physiology and immune defense (Vázquez and Guerrero, 2007). Nonspecific and specific immunity in fish has been positively correlated with the number of red and white blood cells. Therefore, total blood cell count may be a convenient indicator of fish physiological status. Damage to blood organs in fish suppress immune ability and even endanger organism survival. *A. hydrophila* infection decreased the blood cell numbers and modulated the innate immune factors in olive barb (Das et al., 2011).

Reactive oxygen species (ROS) are commonly used as messenger molecules in normal cell functions. However, overproduction of ROS may lead to oxidative damage to tissue macromolecules including DNA, proteins, and lipids (Wang et al., 2009), and subsequently induce cell apoptosis (Circu and Aw, 2010). Calcium has been recognized as an important secondary messenger for a variety of cellular processes. Oxidative stress causes a Ca^{2+} influx into the cytoplasm from the extra-cellular environment, and the endoplasmic reticulum or sarcoplasmic reticulum (ER/SR) through the cell membrane, and the ER/SR channels, respectively (Wang et al., 2009). Ca^{2+} overload or disturbances in local intracellular distribution lead to cell death (Orrenius et al., 2003). Nitric oxide (NO) has been identified as a vital physiological modulator and signaling molecule in both mammals and fish (Acosta et al., 2004). The dysregulation or overproduction of NO has been involved in immune system damage and cell apoptosis (Kim et al., 2001).

River pufferfish *Takifugu obscurus*, widely distributed in the Sea of Japan, the East China Sea, and the Yellow Sea, is an anadromous fish. Because of the relatively small size and simple organization of its genome, *Takifugu* is used as a model organism to understand peculiar physiology, morphology, and genomics (Aparicio et al., 2002). In recent years, farmed pufferfish have become increasingly popular in Chinese markets due to their rapid growth and high

nutritional quality. However, pufferfish culture suffers from serious disease problems caused by bacterial pathogens. In this study, we investigated the effects of *A. hydrophila* infection on total blood cell count, ROS production, cytoplasmic free- Ca^{2+} concentration, NO production, apoptotic cell ratio, and the transcription levels of caspase-3 in pufferfish. This study can help to further understand the mechanism of cell apoptosis in response to bacterial infection.

Materials and methods

Pufferfish (average body weight 11.80 ± 0.43 g) were supplied by a fish farm in Panyu (Guangdong, China). The fish were kept in 250L cycling-filtered plastic tanks containing continuously circulating aerated water at $25 \pm 1^\circ\text{C}$ (pH 7.5; 6.2 mg/L dissolved oxygen) for two weeks before experimental treatments. During the acclimation period, the commercial fish diet (42% protein, 8.0% fat, 5.0% fiber and 15% ash, supplied by a commercial diet, China) was fed twice a day at a rate of 3% wet body weight.

Aeromonas hydrophila (*A. hydrophila*) was obtained from diseased pufferfish. The isolate was confirmed as *A. hydrophila* after biochemical and molecular identification. *A. hydrophila* was cultured in Lysogeny broth (LB medium) at 28°C for 12 h. The bacterial cells were collected by centrifugation (3000 rpm, 10 min) and washed twice with phosphate-buffered saline (PBS). The bacteria were then resuspended in PBS. Fish were infected with a 50% 72-h LD_{50} dose of bacteria (1×10^7 CFU/ml). One hundred and sixty fish were randomly divided into two groups (80 individuals/group) the control group, and the treatment group. Each fish in the treatment group was injected intraperitoneally with 100 μL of suspended bacteria, while the control group was injected with the same volume of PBS. At 0, 3, 6, 12, 24, 48, and 72 h, six fish from each group were randomly sampled and dissected after anesthesia in 0.05% tricaine methane sulfonate (MS-222, Sigma Diagnostics INS, St. Louis, MO) respectively. Blood samples were taken from the heart using a sterilized syringe containing an anticoagulant solution and immediately transferred into an individual tube held on ice to measure total blood cell count, oxygen species (ROS) production, cytoplasmic free- Ca^{2+} (cf- Ca^{2+}) concentration, intracellular nitric oxide (NO) production, and apoptotic cell ratio (see below).

A drop of diluted blood sample (10 μL) was placed on a hemocytometer and counted under a light microscope (Olympus).

To monitor the level of respiratory bursts, the cell-permeant probe 2',7'-dichlorofluorescein diacetate (DCFH-DA, Sigma) was used as described by Xian et al. (2009). A volume of 200 μL blood cells suspension was incubated with 10 μM DCFH-DA for 30 min in the dark at room temperature. Then the fluorescence of the cell suspensions was analyzed using the flow cytometer (Becton-Dickinson FACSCalibur). Typically, 10,000 cells were analyzed for the two fluorescent signals. ROS production was expressed as mean fluorescence of DCF.

The cf- Ca^{2+} concentration in the current experiment was examined using the previously described method (Xian et al., 2010). A volume of 200 μL blood cells suspension was incubated with 10 μM fluo-3/AM for 30 min in the dark. Then the fluorescence of the cell suspensions was analyzed by flow cytometer.

The fluorescent probe 4-amino-5-methylamino-2',7'-difluorofluorescein diacetate (DAF-FM DA, Sigma) was used to measure intracellular NO production. DAF-FM DA solutions were prepared by dissolving DAF-FM DA in dimethyl sulphoxide (DMSO, Sigma Diagnostics INS, St. Louis, MO) to 1 mM and stored at 20°C . 200 μL blood cell suspension was incubated with 10 μM DAF-FM DA for 60 min at room temperature in the dark, then the DAF-FM fluorescence of cells was recorded on FL1 by detector of flow cytometer. NO production was expressed as mean fluorescence of DAF-FM in arbitrary units (A.U.).

The apoptosis of blood cells were determined with Annexin V-FITC and propidium iodide (PI) staining using an apoptosis detection kit (Invitrogen)

following the manufacturer's instructions. The collected blood cells were diluted with anticoagulant solution to obtain a final concentration of 1×10^6 cells. The pellets were resuspended by Annexin V-FITC binding buffer, and then incubated with Annexin V-FITC in the dark at 20-25°C for 10 min. Samples were then analyzed by a flow cytometer.

Total RNA was extracted using TRIzol reagent (Invitrogen, USA) according to the manufacturer's instructions. The A260/280 ratios of all the RNAs prepared were measured by Nanodrop 2000 spectrophotometer (Nanodrop Technologies, Wilmington, DE, USA). RNA integrity was checked by electrophoresis on 1% agarose gels. Single-stranded cDNA was synthesized from 1 µg total RNA using PrimeScript RT reagent Kit With gDNA Eraser (Takara, Dalian, China) following the manufacturer's instructions. cDNA templates were then stored at -80°C for later analysis.

mRNA expression levels of caspase-3 in blood cells after bacterial inoculation were measured by quantitative real time RT-PCR. Information of gene specific primers used for real-time PCR is given in Table 1. Pufferfish beta actin was selected as a house keeping gene in this study. Before the qRT-PCR experiments, the specificity and efficiency of the primers above were detected. Real-time PCR was amplified in an ABI 7500 real-time PCR machine (Applied Biosystems, USA) using SYBR Premix Ex Taq™ (Takara, Dalian, China) following the manufacturer's recommendations. Reaction mixtures were 20 µL, containing 2 µL cDNA sample, 0.4 µL ROX, 10 µL 2× SYBR Premix Ex Taq, 0.4 µL each of the 10 mM forward and reverse primers, and 6.8 µL dH₂O. The real-time PCR conditions were as follows: 94°C for 10 min, then 45 cycles at 95°C for 30 s, 60°C for 30 s and 72°C for 30 s, followed by 10 min at 72°C. After the program was completed, the threshold cycle (Ct) values were obtained from each sample. Relative gene expression levels were evaluated using $2^{-\Delta\Delta C_T}$ method (Livak and Schmittgen, 2001)

Table 1. The sequences of primers in this experiment.

Target gene	Primer sequence (5'-3')	Gen Bank	Amplicon size (pb)
Caspase 3	F:CGAGGGCGTGTTCCTTGGT R:GGGATCTTGGTGGTGCTGC	NM_001032699.1	135
β-Actin	F:CATCACCATCGGCAACGAGAGG R:CGTCGCACTTCATGATGCTGTG	AJ715484.1	115

All data are expressed as means ± standard deviation. Significant differences were evaluated by a one-way ANOVA followed by Duncan's multiple range tests. Statistical analysis was performed using SPSS 18.0 software (SPSS, Chicago, IL, USA). *P* value < 0.05 was considered to be statistically significant.

Results

Blood cell numbers after bacterial infection are shown in Fig. 1. At the beginning, the mean blood cell count was 122.7×10^7 cells/ml. After bacterial infection, the blood cell count decreased significantly at 3, 6, 12, 24, 48 and 72 h. The lowest level of blood cell count was observed at 48 h post-infection.

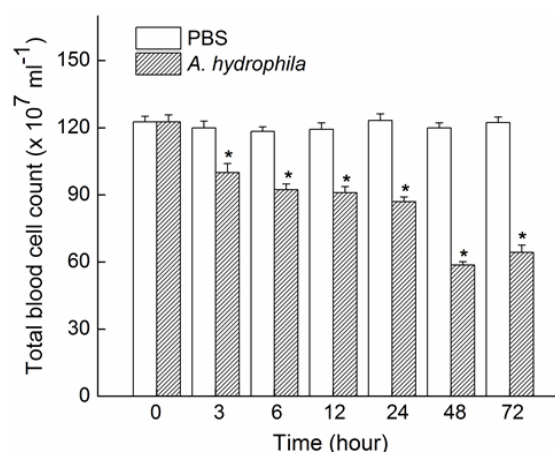


Fig.1. Total blood cell count of pufferfish in response to *A. hydrophila* infection.

ROS levels in blood cells are shown in Fig. 2. ROS production increased significantly at 3 h after bacterial infection ($P < 0.05$). Moreover, the highest ROS levels were observed at 72 h post-infection.

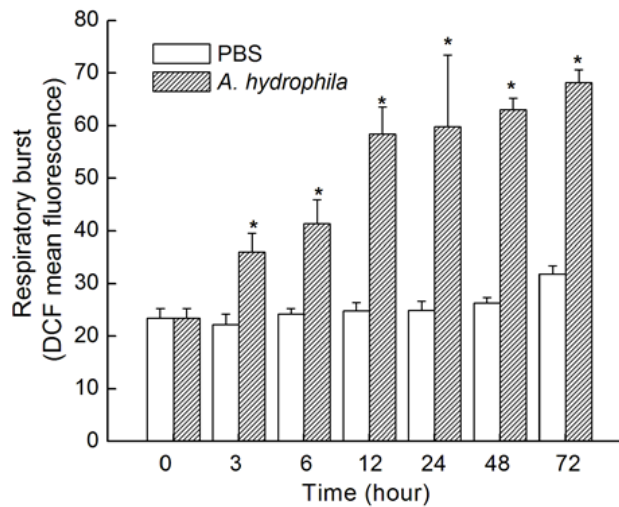


Fig.2. Respiratory burst activity in the blood cells of pufferfish in response to *A. hydrophila* infection.

As shown in Fig. 3, there were no significant differences in cf-Ca^{2+} concentration at 3 h and 6 h post-infection.

Cf-Ca^{2+} concentration increased significantly at 12 h, and then increased further to a high level at 48 h, but then decreased (more than the control) until the end of the experiment.

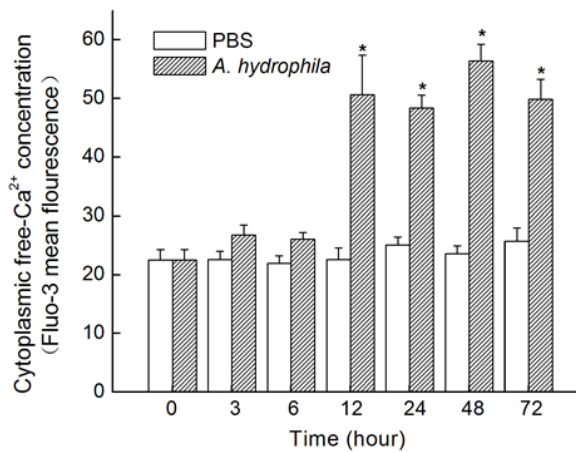


Fig.3. Nitric oxide production in the blood cells of pufferfish in response to *A. hydrophila* infection.

The changes of the NO production after bacterial infection were shown in Fig. 4. NO production increased significantly at 3, 6, 12, 24, 48 and 72 h after bacterial infection. The highest production of NO was observed at 72 h post-infection.

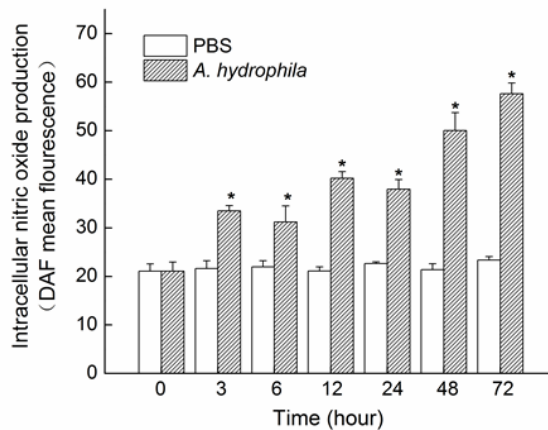


Fig.4. Cytoplasmic free- Ca^{2+} concentration in the blood cells of pufferfish in response to *A. hydrophila* injection.

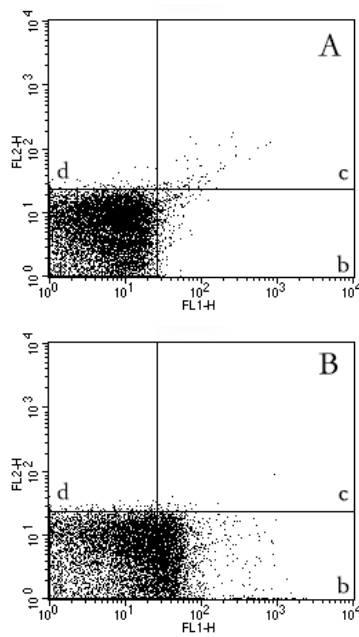


Fig.5. Apoptosis dot plot of blood cells of pufferfish after 0 h (A) and 72 h (B) *A. hydrophila* infection. Live cells are located in quadrant (a). Early apoptotic cells are located in quadrant (b). Late apoptotic cells and necrotic cells are located in quadrant (c). Cell debris is located in quadrant (d).

At the beginning, the percentage of apoptotic cell was 1.91% (Fig. 5, 6). Apoptotic cell ratio of blood cells increased significantly to 10.82%, 12.84%, 25.85%, and 27.74% at 3, 6, 12 and 24 h post-infection, respectively. The highest percentage of apoptotic cell (63.92%) was observed at 48 h. Then, percentage of apoptotic cell decreased to 38.21% at 72 h.

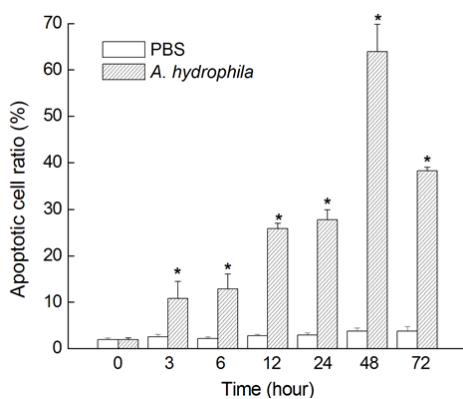


Fig.6. Percentage of apoptotic cell in the blood cells of pufferfish in response to *A. hydrophila* infection.

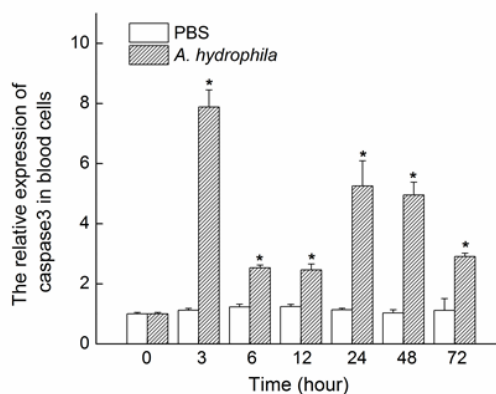


Fig.7. Expression of the caspase-3 transcript at 0, 3, 6, 12, 24, 48 and 72 h after *A. hydrophila* injection in the blood cells.

The change of caspase-3 transcription was investigated after inoculation with bacteria. As shown in Fig. 7, the caspase-3 transcription levels were significantly up-regulated after bacterial infection. The highest expression fold in response to bacterial challenge occurred at 3 h (7.9-fold).

Discussion

Aeromonas hydrophila is distributed widely in aquatic environments and can infect terrestrial animals and a variety of farmed fish. It is a causative agent of motile aeromonad septicemia. Disease outbreaks usually occur when fish are immune-compromised by stresses from overcrowding, low oxygen, high temperature or concurrent diseases (Stevenson, 1988; Shao et al., 2004). Previous study demonstrated that *A. hydrophila* induced apoptosis in fish as well as in

mammalian cells (Shao et al., 2004). Little is known about the relationship between cell apoptosis and pathogenesis of fish bacterial diseases. In the present study, we investigated the effects of *A. hydrophila* infection on total blood cell count, oxygen species (ROS) production, cytoplasmic free- Ca^{2+} (cf- Ca^{2+}) concentration, intracellular nitric oxide (NO) production, apoptotic cell ratio, and the transcription levels of caspase-3 in pufferfish.

Blood is an important indicator for immune function and oxygen transportation (Vázquez and Guerrero, 2007; Singh and Srivastava, 2010). Decrease in blood cell numbers might lead to impaired immune capability, decreased resistance to bacterial infection, and ultimately death (Li et al., 2014). *A. hydrophila* infection has been reported to result in decreased blood cell count in fish (Das et al, 2001). In this study, a sharp decrease of the blood cell count was observed after *A. hydrophila* infection. Decreased blood cell count may lead to impaired immune capability and ultimately death.

The release of ROS by blood cells is a critical step in the innate immune response by which potential pathogens and parasites are eliminated following phagocytosis (Xian et al, 2010). Under normal conditions, mild increase in ROS is considered to be beneficial with respect to increased immunity. However, high ROS such as hydrogen peroxide (H_2O_2), free radicals superoxide ($\text{O}_2^{\bullet-}$) and hydroxyl radical (HO^{\bullet}) can be important mediators of damage to the host cell (Qi, et al, 2013). Previous study has showed that infections and diseases increase the production of ROS which may damage important biomolecules, such as DNA, proteins and lipids (Halliwell, 1993). Moreover, ROS is an important factor in apoptotic processes and acts as key agents in the events leading to neurodegeneration, such as caspase activation, glial activation, mitochondrial dysfunction, protein misfolding, and proteasomal dysregulation (Andersen, 2004; Wang, et al, 2008). In our study, ROS level of blood cells after bacterial infection was significantly higher than that of the control group. This result suggested that bacterial infection could induce excessive ROS production.

As an important second messenger, calcium signaling is known to be involved in regulating a variety of cellular processes. Ca^{2+} signaling may also modulate many physiological or pathological processes including apoptosis (Xian, et al, 2013). Some previous studies have shown that an increase in cytosolic Ca^{2+} may lead to apoptosis due to stimulation of intracellular protease(s), nucleases, phospholipases, and other hydrolytic enzyme activities (Orrenius, 2003; Chen, et al, 2006). It has been shown that an increase of cf- Ca^{2+} concentration may be stimulated by ROS (Rizzuto et al., 2003). In this study, cf- Ca^{2+} concentration increased significantly at 12 h after bacterial infection. The possible role of calcium in *A. hydrophila*-induced apoptosis has also been reported in fish cells (Banerjee et al., 2012). These alterations in cf- Ca^{2+} concentration were consistent with elevated production of ROS. This result suggested that cf- Ca^{2+} may be an apoptotic signal mediator, playing a major role in the mechanism of programmed death induced by bacterial infection.

NO is a multifunctional free radical generated by three different forms of nitric oxide synthases (NOS): inducible NOS (iNOS), neuronal NOS (nNOS), and endothelial NOS (eNOS) (Pavanelli et al., 2010; Jin et al., 2011). It has been reported that NO is produced via the induction of iNOS by macrophages in response to outside stimuli (Tafalla et al., 2005). Previous studies suggested that the overproduction of NO was involved in cell apoptosis (Kim et al., 2001; Lu et al., 2003). To initiate apoptosis, NO can induce mitochondrial dysfunction by stimulating membrane permeability transition thus activating a cascade of reactions leading to cell death (Marriott et al., 2004; Wang et al., 2008). In our study, we observed that NO production increased significantly 3h after bacterial infection. This result suggested that bacterial infection could change NO production.

Mitochondria are one of the most important organelles in apoptosis (Christophe et al., 2002). Mitochondrial alterations such as loss of transmembrane potential, production of ROS, disruption of the electron transport chain, and decrease in ATP synthesis are tightly related to apoptosis (Xiang et al., 2008). Activation of the apoptosis process can enhance immune defense by limiting the spread of pathogens, and preventing inflammatory damage of surrounding tissues (Mai et al., 2010). In this study, the apoptotic blood cells were examined using flow cytometer. Apoptotic cell ratio increased significantly after *A. hydrophila* infection. This result had a high correlation with increased ROS, cf-Ca²⁺, and NO production. Therefore, our results further suggested that bacterial infection could cause a variety of disorders in the blood system and trigger apoptotic and necrotic cell death.

Apoptosis is modulated by the extrinsic or intrinsic apoptotic pathways that involve a series of biochemical regulators and molecular interactions (Bridgham et al., 2003; Wang et al., 2008). The central molecules of apoptosis are the caspases. Both extrinsic and intrinsic apoptotic pathways result in the activation of caspase-3, which is a major executioner of apoptosis in the apoptotic pathway (Li et al., 2011). Activation of caspase-3 causes many of the characteristic morphological alterations of apoptosis, such as breakdown of several cytoskeletal proteins, and degradation of the inhibitor of caspase-activated DNases, directing the cell toward death (Cohen 1997; Elvitigala et al., 2012). In the present study, we observed the ascension caspase-3 activity after *A. hydrophila* inoculation, suggesting that *A. hydrophila* could induce caspase-dependent apoptosis in pufferfish blood cells.

In conclusion, the present study demonstrated that *A. hydrophila* infection altered ROS, NO and cf-Ca²⁺ levels, and activated caspase-3, which then led to cell apoptosis. The information presented in this study will help to elucidate the precise mechanism of cell apoptosis in response to bacterial infection.

Acknowledgements

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