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# PI3K signaling pathways modulated white spot syndrome virus (WSSV) replication in *Procambarus clarkii*



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### ABSTRACT

The PI3K/AKT signaling pathway is commonly exploited to regulate viral replication and affect the fate of infected cells. In the present study, a PI3K-specific inhibitor (LY294002) was employed to pretreat crayfish to evaluate the effects of PI3K/AKT signaling pathway in WSSV replication. The results showed that the WSSV copy numbers in crayfish pretreated with LY294002 were significantly lower than those in Tris-HCl pretreatment crayfish on the sixth and tenth day after WSSV infection. In semigranular cells, the apoptosis rates were upregulated on the third day post-WSSV infection, and a significantly lower proportion of apoptosis cells were observed in LY294002-pretreatment group. The expression level of Bax, Bax inhibitor-1 and lectin mRNA in haemocytes of crayfish were increased after WSSV infection. After the secondary stimulation with Tris-HCl, the Bax expression level in LY294002-pretreatment crayfish was significantly higher than that of crayfish pretreated with Tris-HCl on the third or sixth day, but the Toll and lectin mRNA expression decreased significantly on the third, sixth and tenth day. The Bax mRNA expression levels in LY294002-WSSV group were significantly higher than those in Tris-HCl-WSSV group on the third and tenth day. The Bax inhibitor-1 mRNA expression levels in LY294002-WSSV group were significantly lower than those in Tris-HCl-WSSV crayfish on the third day. These results together indicated that the hosts PI3K/AKT signaling pathway play positive roles in WSSV replication through the balance between host cell apoptois and innate immune responses. This information is helpful to further understand the role of PI3K/AKT signaling pathway on WSSV replication in Decapoda crustaceans.

### 1. Introduction

The phosphatidylinositol-kinase (PI3K)-AKT signaling pathway regulates several key cellular functions, such as protein synthesis, glucose metabolism, cell growth and survival [1]. It is established that viral infections exploit PI3K/AKT signaling pathway to regulate viral cellular entry, ensure successful virus replication, drive viral protein expression, as well as the suppression of the host's antiviral activities [2]. Virus modulate PI3K/AKT signaling pathway to enter host cells, such as Zaire Ebola virus (ZEV), human immunodeficiency virus type 1 (HIV-1), Avian leukosis virus (ALV), human herpes viruses (HHV), cytomegalovirus (CMV) and herpes simplex virus (HSV) [3–7]. The PI3K/AKT signaling pathway has been demonstrated to be required not only for viral cell entry but also for subsequent viral replication [6]. For

example, inhibition of PI3K activation by a PI3K-specific inhibitor (LY294002) during porcine circovirus type 2 (PVC2) and respiratory syndrome virus (PRRSV) infection both led to lower virus yield, which is associated with decreased viral DNA replication and lower virus protein expression [8,9]. Influenza A virus also activated the PI3K/AKT signaling pathway to augment its efficient replication [10]. Therefore, to complete the viral life cycle, PI3K/AKT signaling pathway is commonly exploited to regulate viral replication and affect the fate of infected cells.

Apoptosis is defined as an active physiological process of cellular self-destruction, which is also utilized for viruses to evade the host immune system to ensure viral replication, viral spread, persistent infection and inflammatory responses [11]. The process of apoptosis is controlled by different ranges of cellular signaling pathways. Inhibition

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of PI3K activation with LY294002 could reduce HSV protein expression and increase the cleavage and release of caspase-3 [6]. Likewise, during largemouth Bass virus (LMBV), dengue virus (DEN) or Japanese encephalitis virus (JEV) infection, blockage of PI3K activation by LY294002 greatly enhanced virus-induced apoptosis at an early stage of infection [11–13]. Leu et al. proposed that invasion of white spot syndrome virus (WSSV) could lead to increase expression of proapoptosis proteins, membrane permeabilization of mitochondria, and cellular apoptosis [14]. Although the apoptosis has been documented to occur in several shrimps infected with WSSV, such as *Litopenaeus vannamei* and *Penaeus monodon* [15,16], the PI3K/AKT signaling pathway involved in these processes still remained largely unknown.

WSSV has a broad host range of Decapoda crustaceans, such as penaeid shrimp, crab and crayfish. It is difficult to prevent and inhibit the spread of the virus due to the wide host range and easy replication, which caused huge economic losses during the last three decades [17]. However, Invertebrates have developed various defense mechanisms to recognize and eliminate the potential pathogens during the process of evolution. Recognition of non-self is the first line in mounting immune responses, and it is mediated by a group of proteins named pattern recognition receptors (PRRs). There are growing evidence that PRRs (i.e., Toll-like receptor and lectin) are closely related to virus replications in hosts [18,19]. Toll receptors are evolutionary ancient families of PRRs with crucial roles in invertebrate innate immune response against WSSV [20]. And lectins are one type of important PRRs that widely distribute in almost all organisms and involve in many biological processes, including cell-cell interaction, protein synthesis, and signal transduction [21]. Although PRRs involved in host immune defense responses against WSSV have been characterized, the potential biological mechanism is currently unclear. The studies of WSSV replication and host-interactions would contribute to the development management strategies for WSSV prevention and disease control.

The freshwater crayfish, *Procambarus clarkii*, is one of the most important commercially cultivated crustaceans in China. Based on the previous study of enrichment of the PI3K/AKT signal pathway in the regulation of viral replication, we investigated the role and mechanisms of PI3K/AKT regulation during WSSV infection in the crayfish *P. clarkii*. In the present study, a PI3K-specific inhibitor (LY294002) was employed to pretreat crayfish comparatively to evaluate the effects of PI3K/AKT signaling pathway in WSSV replication. Alteration of WSSV copy numbers, cell apoptosis, the apoptotic associated factors (Bax and Bax inhibitor-1) and PRRs (Toll-like receptor and lectin) were measured to elucidate the potential mechanism of PI3K/AKT signaling pathway in the crayfish *P. clarkii* against WSSV. Taken together, these results of the interaction between WSSV replication and the PI3K/AKT signaling pathway would provide new insight into the mechanism of WSSV pathogenesis in crayfish or other commercial Decapoda crustaceans.

### 2. Materials and methods

### 2.1. Crayfish and virus preparation

Crayfish *P. clarkii* (approximately  $21.3 \pm 1.8$  g each) were separately collected from aquatic markets in Yantai, Shandong Province, Weishan Lake, Shandong Province and Lianyungang, Jiangsu Province, China, and cultured in aerated water tanks with fresh water at  $25 \pm 2$ °C for two weeks before processing.

The shrimps with WSSV were provided from Yellow Sea Fisheries Research, Institute Chinese Academy of Fishery Sciences. WSSV was purified by differential centrifugation in shrimps. Briefly, 1 g of the gill tissue from the infected shrimps was homogenized in 45 mL of TNE buffer (50 mM Tris-HCl, pH7.5, 400 mM NaCl, 5 mM EDTA, pH 8.5) with 1 mM PMSF. Thereafter, it was centrifuged at  $3500 \times g$  for 5 min at 4 °C, and the supernatant was then centrifuged at  $30000 \times g$ for 30 min at 4 °C. The precipitate was resuspended in 10 mL TM buffer (50 mM Tris-HCl, 10 mM MgCl<sub>2</sub>, pH7.5) and then centrifuged at

### Table 1

Names and sequences of primers used in this study.

Primer name	Sequence
WSSV-F154 WSSV-R154 TaqMan probe	CCAGTTCAGAATCGGACGTT AAAGACGCCTACCCTGTTGA TCCATAGTTCCTGGTTTGTAATGTGCCG
Oligo (dT)-adaptor	GGCCACGCGTCGACTAGTAC(T)17
Bax Pc-Bax-RT-F Pc-Bax-RT-R	TATAGTTGGCTCATTAGCAG ATACTAAGTGAAGATGACTG
Bax inhibitor-1 PcBI-1-RT-F PcBI-1-RT-R	TGCCATTACATCTTGGGTTCT CGACCTAATCCCATCTCAAGC
lectin RT-PcL-lecF RT-PcL-lecR	CTTGGTGGCTGCCTTAGAG TCCTTCTCATTGTTTGTCATTC
Toll Toll-F Toll-R	TCATTTGGCATCTGGCTCAC GCAGGTGGTGGCGTTGA
18sRNA 18s RNA-RT-F 18s RNA-RT-R	TCTTCTTAGAGGGATTAGCGG AAGGGGATTGAACGGGTTA

 $30000 \times g$  for 20 min at 4 °C. Finally, the precipitate was resuspended in 1 mL TM buffer with 0.3%  $\rm NaN_3$  as WSSV stock solution and stored at -80 °C until use.

### 2.2. WSSV quantification

The standard curve for WSSV quantification was performed according to a TaqMan real-time assay [17]. A WSSV DNA fragment of 154 bp amplified using primers WSSV-F154 and WSSV-R154 (Table 1) for real-time PCR was ligated into pMD18T-Easy vector and cloned into DH5 $\alpha$ . The WSSV copy numbers of the target amplicon in the plasmid were estimated, and target amplicon in the plasmid was used to make 10-fold serial dilutions. The TaqMan real-time PCR was assayed in a PCR mixture containing 0.25  $\mu$ M WSSV-F154, 0.25  $\mu$ M WSSV-R154, 0.125  $\mu$ M TaqMan probe (Table 1) and the serial dilutions of genomic DNA in a final reaction volume of 20  $\mu$ L. Regression of the log of WSSV copy number and the corresponding CT value was used as a standard curve for determining viral load. The viral copy number was normalized on a ng genomic DNA basis.

### 2.3. Drug treatment

A PI3K-specific inhibitors LY294002 was purchased from Cell Signaling Technology, which was dissolved in 10% DMSO and diluted with Tris-HCl (50 mM, pH7.5). Thirty crayfish were used to survey the phosphorylated AKT expression level after LY294002 treatment. Ten crayfish were injected with LY294002 (0.625  $\mu$ g·g<sup>-1</sup>, 100  $\mu$ L) or Tris-HCl (100  $\mu$ L) respectively, and the untreated crayfish were employed as the blank group. Five crayfish were randomly sampled for western blot analysis at 2 h post challenge.

### 2.4. Western blot analysis

The haemocytes in crayfish were lysed with 1 mL cell lysis buffer (50 mM Tris-HCl, 150 mM NaCl, 1.0% Triton X-100, 1.0% sodium deoxycholate, and 0.1% sodium dodecyl sulfate [SDS]). The protein content was measured using the BCA Protein Assay Kit (Beyotime, Shanghai, China) and 150 mg total protein was subjected to 12% sodium dodecyl sulfate polyacrylamide gel electrophoresis and trans-ferred to polyvinylidene difluoride membranes. The membranes were incubated with an anti-phospho-AKT (pAKT) rabbit hAb or an anti-AKT rabbit hAb (Cell Signaling Technology, Inc, USA) overnight at 4.0 °C, followed by incubation with anti-rabbit IgG HRP-linked antibody (Cell Signaling Technology, Inc, USA) for 1.0 h at room temperature. The blots were detected with an enhanced chemiluminescence detection kit according to the manufacturer's instruction (Beyotime, Shanghai, China).

### 2.5. WSSV infection

Two hundred and eighty crayfish were used in the experiment, and they were divided into seven groups. In the first group, forty crayfish were untreated and employed as blank group. In group II or III, Forty cravfish were treated with Tris-HCl (pH7.5, 100 µL) or WSSV stock (10<sup>7</sup>  $copies uL^{-1}$ , 100 uL), respectively. In group IV, the forty cravfish were injected with Tris-HCl (100 µL) twice with 2 h interval. In group V, the forty crayfish were pretreated with LY294002 ( $0.625 \,\mu g \, g^{-1}$ ,100  $\mu L$ ) firstly and then injected with Tris-HCl (100 µL) at 2 h after the first stimulation. In group VI, the crayfish were injected with Tris-HCl (100  $\mu L$ ), and then received an injection of WSSV stock (10  $^7$ copies  $\mu L^{-1}$ , 100  $\mu L$ ) at 2 h after the first stimulation. In group VII, the forty crayfish were pretreated with LY294002 (0.625  $\mu g \cdot g^{-1}$ ,100  $\mu L$ ), and then they were received a secondary injection of WSSV stock (10<sup>7</sup> copies  $\mu$ L<sup>-1</sup>, 100  $\mu$ L) at 2 h after the fist stimulation. Afterwards, the crayfish in every group were divided into two shares, and the crayfish of one share were used to monitor the numbers of dead crayfish and the biological issue when they got moribund. In the other share, five crayfish of each group were randomly sampled on the third, sixth and tenth day after treatment to detect alteration of WSSV copy numbers, cell apoptosis rates, and the associated factors expression (Bax, Bax inhibitor-1, Toll and lectin).

### 2.6. DNA extraction and real-time PCR amplification

Total genomic DNA was extracted from crayfish treated with WSSV by using a tissue DNA Kit (OMEGA, USA) for detecting the WSSV copy numbers. The DNA concentrations were determined by optical density at OD 260 nm using a Genesys 5 spectrophotometer (Spectronic instrument, Inc., New York, USA). Real-time PCR was performed by using a SYBR Premix Ex TaqTM (RR820A, Takara) containing a high-performance Taq antibody and Takara Ex Taq HS. A sample of total DNA was added to a PCR mixture containing 0.25  $\mu$ M WSSV-F154, 0.25  $\mu$ M WSSV-R154 and 0.125  $\mu$ M TaqMan probe in a final reaction volume of 20  $\mu$ L. The amplification program consisted of 30 s at 95 °C for initial denaturation followed by 45 cycles of 5 s at 95 °C for denaturation and annealing at 35 s at 60 °C. The WSSV copy numbers were calculated according to the method mentioned above.

### 2.7. Haemocytes collection

From each crayfish, about 0.5 mL hemolymph was collected into the tube containing equal volume of anticoagulant solution (100 mM glucose, 140 mM NaCl, 30 mM trisodium citrate, 26 mM citric acid, 10 mM EDTA·Na<sub>2</sub>·2H<sub>2</sub>O, pH 4.6). The hemolymph was collected and centrifuged at 800 × g, 4 °C for 10 min to collect the haemocytes. The haemocytes were divided into two shares, one of which was subjected to the measurement of cell apoptosis, and the other one was immediately used for total RNA extraction using Trizol reagent according to the manufacture's protocol (Invitrogen, USA).

### 2.8. Cell apoptosis assays

Cell apoptosis was assessed using an Annexin V-FITC Apoptosis Detection Kit (Becton Dickinson, Franklin Lakes, NJ, USA) according to the manufacturer's instructions [22]. Briefly,  $1.0 \times 10^5$  haemocytes were mixed with 500 µL Annexin Binding Buffer,  $5.0 \,\mu\text{L}$  FITC-conjugated Annexin V antibody, and  $5.0 \,\mu\text{L}$  propidium iodide. The mixture was incubated in the dark for 15 min at room temperature. The relative

number of apoptotic cells was determined using a Beckman Coulter EPICX-XL Flow Cytometer (Beckman Coulter, Inc., Brea, CA, USA). The normal cells without induction of apoptosis were used for fluorescence compensation regulation to remove the spectral overlap.

Crayfish haemocytes were divided into three sub-populations, hyaline cells (HC), semigranular cells (SGC) and granular cells (GC), according to their discrepant values in forward scatter (FSC) and side scatters (SSC) [23]. Flow cytometry analysis of total haemocytes, HC, SGC, and GC in crayfish at different time points was performed to study apoptosis.

### 2.9. Mitochondrial membrane potential ( $\Delta \psi$ )

Mitochondrial membrane potential ( $\Delta \psi$ ) was detected using the BD<sup>TM</sup> MitoScreen Kit (BD Biosciences, USA) according to the manufacturer's instructions. Briefly,  $1.0 \times 10^6$  haemocytes were mixed with 12.375 mL 1 × Assay Buffer, 0.125 mL JC-1 Stock Solution, and 0.5 mL JC-1 Working Solution. The mixture was incubated in the dark for 15 min at 37 °C in CO<sub>2</sub> incubator. The  $\Delta \psi$  evaluated by the JC-1 polymer/monomer fluorescence ratio was determined using a BD FACSAria<sup>TM</sup> III sorter (BD Biosciences, USA).

## 2.10. Quantification of Bax, Bax inhibitor-1, Toll and lectin expression by quantitative real-time RT-PCR

Quantitative real-time polymerase chain reaction (RT-PCR) was performed to investigate Bax, Bax inhibitor-1, Toll and lectin mRNA expression of crayfish. Total RNA was extracted from haemocytes as described above. The SYBR Green RT-PCR assay was performed in an ABI PRISM 7500 Sequence Detection System (ThermoFisher Scientific, USA). The amplifications were conducted in triplicate in 20 µL using the following thermal profile: 95 °C for 3 min, followed by 40 cycles at 95 °C for 15 s and 58 °C for 33 s. The Bax-specific primers Pc-Bax-RT-F and Pc-Bax-RT-R, Bax inhibitor-1-specific primers PcBI-1-RT-F and PcBI-1-RT-R, Toll-specific primers Toll3-F and Toll3-R, lectin-specific primers RT-PcL-lecF and RT-PcL-lecR, and 18s-specific primers 18s RNA-RT-F and 18s RNA-RT-R (Table 1), were used to amplify the corresponding products. The 18s RNA was chosen as reference gene for internal standardization. The comparative average-cycle threshold method was used to analyze the Bax, Bax inhibitor-1, Toll and lectin mRNA expression level. The expression level of the Bax, Bax inhibitor-1, Toll and lectin gene could be calculated, and the value of which stood for an n-fold difference relative to the calibrator [24].

### 2.11. Statistical analysis

Data are presented as mean  $\pm$  standard deviations (SD) from at least five independent experiments. All data were subjected to one-way analysis of variance.  $P \leq 0.05$  was considered statistically significant.

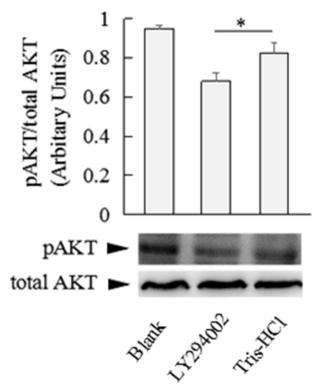
### 3. Results

### 3.1. The expression of phosphorylated AKT

To examine the effect of a PI3K inhibitor (LY294002) on inhibiting the activity of PI3K, the expression level of phosphorylated AKT was tested through western blot analysis. The PI3K inhibitor (LY294002) was able to reduce basal AKT phosphorylation (Fig. 1). At 2 h after LY294002 injection, phosphorylated/total AKT was remarkably decreased and 4/5-fold (P < 0.05) lower than that of Tris-HCl-treatment crayfish.

### 3.2. Crayfish survival rate, standard curve and WSSV quantification

The survival rates of crayfish were equal or greater than 80% on the tenth day and no significant difference among the seven groups was

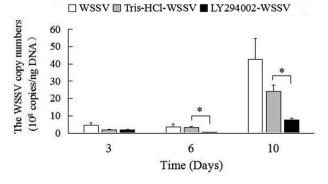


**Fig. 1.** Effects of a PI3K inhibitor (LY294002) on phosphorylation of AKT in crayfish. The crayfish were injected with LY294002 or Tris-HCl respectively, and the untreated crayfish were employed as blank group. Phosphorylated/total AKT were detected using western blot analysis. Each symbol and vertical bars represented the mean  $\pm$  SD (N = 5). Significant differences between LY294002-treatment group and Tris-HCl-treatment group were indicated with one asterisk at P < 0.05.

observed (data not shown). The crayfish in Tris-HCl-WSSV group got moribund on the fifth day after WSSV infection, and the moribund crayfish were observed in the LY294002-pretreatment group on the ninth day after WSSV infection.

Strong linear correlation ( $R^2 = 0.988$ ) was obtained between CT and the WSSV copy numbers ranging from  $1 \times 10^{12}$  to  $1 \times 10^1$  in realtime PCR assay. With an optimal PCR mixture, the genomic DNA of the specimen was detected in one real-time PCR reaction, and the results showed the large dynamic range and high sensitivity of the assay (data not shown).

The WSSV copy numbers in crayfish pretreated with LY294002 were significantly lower than those in Tris-HCl pretreated crayfish on the sixth and tenth day post WSSV infection (Fig. 2). On the third day, the mean WSSV copy numbers in Tris-HCl-WSSV and LY294002-WSSV



**Fig. 2.** The WSSV copy numbers of the crayfish in the three groups (WSSV, Tris-HCl-WSSV and LY294002-WSSV) on the third, sixth and tenth day. Each symbol and vertical bars represented the mean  $\pm$  SD (N = 5). Significant differences between Tris-HCl-WSSV and LY294002-WSSV were indicated with one asterisk at P < 0.05.

group were  $1.78 \times 10^8$  and  $1.83 \times 10^8$  copies ng<sup>-1</sup> DNA, respectively, and no significant difference was observed between Tris-HCl-WSSV and LY294002-WSSV group. On the sixth day, the crayfish in LY294002-WSSV group represented lower WSSV loads with mean values of  $0.59 \times 10^8$  copies ng<sup>-1</sup> DNA, whereas  $3.30 \times 10^8$  copies ng<sup>-1</sup> DNA WSSV copy number was detected in Tris-HCl-WSSV group. The mean WSSV copy number in Tris-HCl-WSSV group was 5.60-fold higher (P < 0.05) than that in LY294002-WSSV group. On the tenth day, the mean WSSV copy number of Tris-HCl-WSSV group was increased and reached 2.41  $\times 10^9$  copies ng<sup>-1</sup> DNA, which was 3.15-fold higher (P < 0.05) than that in LY294002-WSSV group (7.65  $\times 10^8$  copies ng<sup>-1</sup> DNA).

### 3.3. Apoptosis in WSSV-infected haemocytes of crayfish

No substantial changes in apoptosis rates of total haemocytes, HC and GC were observed after infection with WSSV during the whole experiments (data not shown).

According to Annexin V-FITC apoptosis detection method in SGC, there were apoptosis rates of 12.80  $\pm$  3.10% and 9.87  $\pm$  0.68% on the third and sixth day post WSSV infection, which were 1.94-fold (P < 0.05) and 2.69-fold (P < 0.01) higher than those post Tris-HCl stimulation, respectively (Fig. 3). When the crayfish were pretreated with LY294002, a significantly lower proportion of apoptosis cells were observed in Tris-HCl pretreatment group on the sixth and tenth day regardless whether the crayfish were infected with WSSV (Fig. 3). On the sixth day, the apoptosis rate of SGC in LY294002-Tris-HCl group was 6.80  $\pm$  1.00%, which was remarkably lower than that in Tris-HCl-Tris-HCl group (P < 0.05). Meanwhile, 5.40  $\pm$  0.47% of apoptotic rate was obtained in LY294002-WSSV group, and it was also lower significantly than that in Tris-HCl-WSSV group (P < 0.05). On the tenth day, 7.75  $\pm$  0.15% of apoptotic rate was obtained in LY294002-Tris-HCl group. Conversely, apoptotic rate of 13.47 ± 1.64% was found in Tris-HCl-Tris-HCl group. That meant a 1.74-fold (P < 0.05) decrease in LY294002-Tris-HCl group compared to Tris-HCl-Tris-HCl group. In LY294002-WSSV group, 6.17  $\pm$  1.23% of apoptotic rate was detected, and this rate was remarkably lower than that in Tris-HCl-WSSV group (14.77  $\pm$  2.60%) (P < 0.05).

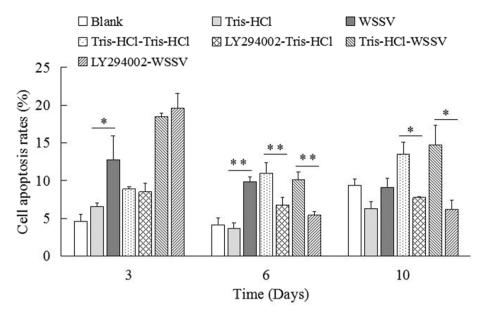
### 3.4. Mitochondrial membrane potential $(\Delta \psi)$ in WSSV-infected haemocytes of crayfish

No substantial changes in  $\Delta \psi$  of total haemocytes, HC and GC were observed after infection with WSSV during the whole experiments (data not shown). In SGC stained with JC-1, there were lower  $\Delta \psi$  (P < 0.01) on the third day post WSSV infection compared to that in crayfish after Tris-HCl injection (Fig. 4). The  $\Delta \psi$  in crayfish pretreated with LY294002 was higher than that in Tris-HCl-pretreatment crayfish on the third day post WSSV infection (P < 0.05) (Fig. 4). On the tenth day, the  $\Delta \psi$  of SGC in LY294002-Tris-HCl group was remarkably higher than that in Tris-HCl-Tris-HCl group (P < 0.05) (Fig. 4).

### 3.5. The Bax mRNA expression in haemocytes of crayfish

The expression level of Bax mRNA in haemocytes of crayfish was increased on the third day after WSSV infection. The relative expression level of Bax transcript in WSSV group on the third day was up-regulated and 5.03-fold higher (P < 0.05) than that in Tris-HCl group (Fig. 5).

There was a significant increase in Bax expression level in LY294002-Tris-HCl group compared to that in Tris-HCl-Tris-HCl group on the third or sixth day (Fig. 5), and the expression levels in LY294002-Tris-HCl group were 11.34-fold (P < 0.05) and 7.77-fold (P < 0.01) higher than that in Tris-HCl-Tris-HCl group, respectively. The expression levels of Bax mRNA in LY294002-WSSV group were significantly higher than those in Tris-HCl-WSSV group on the third and tenth day (P < 0.05) (Fig. 5).



**Fig. 3.** Flow cytometry analysis of apoptosis in semigranular cells (SGC) of seven groups (Blank, Tris-HCl, WSSV, Tris-HCl-Tris-HCl, LY294002-Tris-HCl, Tris-HCl-WSSV and LY294002-WSSV). The rates of apoptotic cells are shown. Vertical bars represent the mean  $\pm$  SD (N = 5). Significant differences between Tris-HCl and WSSV, Tris-HCl-Tris-HCl and LY294002-Tris-HCl, Tris-HCl-WSSV and LY294002-WSSV are indicated with one asterisk at P < 0.05 and two asterisks at P < 0.01.

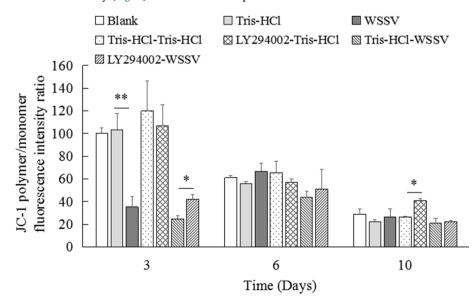
### 3.6. The Bax inhibitor-1 mRNA expression in haemocytes of crayfish

The Bax inhibitor-1 mRNA expression was significantly increased in haemocytes of crayfish after WSSV infection on the third and sixth day (Fig. 6). The relative expression levels of Bax inhibitor-1 in WSSV group on the third and sixth day were up-regulated, and 3.73-fold and 3.95-fold higher (P < 0.05) than that of Tris-HCl group, respectively.

The Bax inhibitor-1 mRNA expression levels in LY294002 pretreatment group were significantly lower than those in Tris-HCl pretreatment crayfish on the third day regardless whether the crayfish were infected with WSSV (Fig. 6). The Bax inhibitor-1 level in LY294002-Tris-HCl group was 0.42-fold lower (P < 0.01) than that in Tris-HCl-Tris-HCl group, and Bax inhibitor-1 level in LY294002-WSSV group was 0.33-fold lower (P < 0.05) than that of Tris-HCl-WSSV group on the third day.

### 3.7. The Toll mRNA expression in haemocytes of crayfish

The Toll mRNA expression was also decreased in LY294002-Tris-HCl group compared to that in Tris-HCl-Tris-HCl group on the third, sixth and tenth day (Fig. 7). The Toll mRNA expression in LY294002-



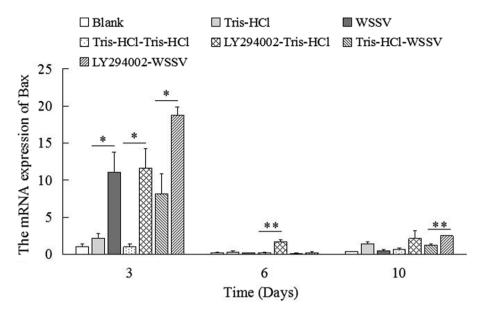
Tris-HCl group decreased significantly than those in the Tris-HCl-Tris-HCl group on the third, sixth and tenth day, respectively (P < 0.05).

### 3.8. The lectin mRNA expression in haemocytes of crayfish

The lectin mRNA expression was significantly increased in haemocytes of crayfish after WSSV infection on the third, sixth and tenth day. When compared to Tris-HCl group, it was 4.42-fold, 2.55-fold and 2.30fold higher in the relative abundance of the lectin mRNA in WSSV group on the third, sixth and tenth day, respectively (P < 0.01) (Fig. 8).

There was a significant decrease in lectin expression level in LY294002-Tris-HCl group compared to Tris-HCl-Tris-HCl group on the sixth and tenth day (Fig. 8). The lectin levels of LY294002-Tris-HCl group were significantly down-regulated on the sixth and tenth day, which were about 0.39-fold (P < 0.05) and 0.19-fold lower (P < 0.01) than those of Tris-HCl-Tris-HCl group on the sixth and tenth day, respectively.

Fig. 4. The  $\Delta \psi$  were evaluated by the JC-1 polymer/ monomer fluorescence intensity ratio in the seven groups (Blank, Tris-HCl, WSSV, Tris-HCl-Tris-HCl, LY294002-Tris-HCl, Tris-HCl-WSSV and LY294002-WSSV) on the third, sixth and tenth day. Each symbol and vertical bars represented the mean  $\pm$  SD (N = 5). Significant differences between Tris-HCl and WSSV, Tris-HCl-Tris-HCl and LY294002-Tris-HCl, Tris-HCl-WSSV and LY294002-WSSV were indicated with one asterisk at P < 0.05 and two asterisks at P < 0.01.



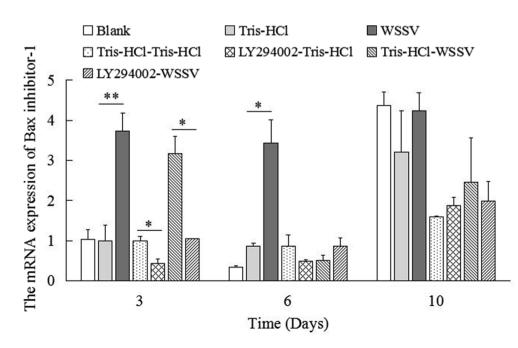
**Fig. 5.** The quantification of Bax transcript relative to 18s RNA transcript in the seven groups (Blank, Tris-HCl, WSSV, Tris-HCl-Tris-HCl, LY294002-Tris-HCl, Tris-HCl-WSSV and LY294002-WSSV) on the third, sixth and tenth day. Each symbol and vertical bars represented the mean  $\pm$  SO (N = 5). Significant differences between Tris-HCl and WSSV, Tris-HCl-Tris-HCl and LY294002-Tris-HCl, Tris-HCl-WSSV ware indicated with one asterisk at P < 0.05 and two asterisks at P < 0.01.

### 4. Discussion

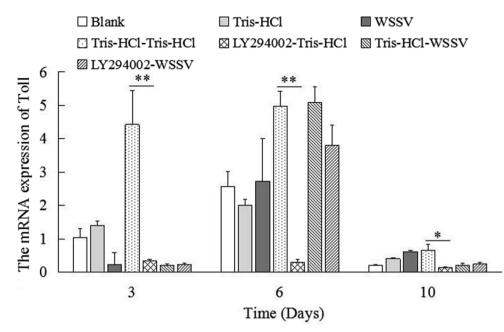
Activation of the PI3K/AKT signaling pathway is common to viral infection, and many virus manipulated this signaling pathway to ensure successful virus replication. The blockage of PI3K/AKT signaling pathway would inhibit viral replication, such as CMV and PRRSV [7,9]. Liu and Cohen demonstrated that inhibition of PI3K/AKT signaling pathway with LY294002 could delay CMV entry, reduce viral immediate-early and early gene expression, and viral DNA replication [25]. Similarly, a significant reduction in PRRSV virus titers was observed in the culture supernatant, when monocytes derived-dendritic cells were pre-treated with LY294002 followed by PRRSV infection [9]. In the present study, the PI3K inhibitor (LY294002) was able to reduce basal AKT phosphorylation (Fig. 1), which showed that this inhibitor was affecting the activity of PI3K in the haemocytes of crayfish. To further test whether PI3K/AKT signaling pathway was involved in WSSV replication, we treated crayfish with LY294002 and detected the WSSV copy numbers after WSSV infection. The results indicated that the WSSV copy numbers in crayfish pretreated with LY294002 were

significantly lower than those in Tris-HCl pretreatment crayfish on the sixth and tenth day post WSSV infection (Fig. 2), which indicated that blockage of the PI3K/AKT signaling pathway could inhibit WSSV replication in crayfish. Viruses are to exploit and manipulate the host pathways to establish an environment favorable for their replication and persistence infection [14]. We demonstrated that the host PI3K/AKT signaling pathway play positive roles in WSSV replication.

During the co-evolution of virus and host, apoptosis was employed by the host as one of the protective measures to prevent and promote virus clearance, which is also utilized for viruses to evade the host immune system to ensure viral replication [26]. In the present study, there were apoptosis rates of 12.80  $\pm$  3.10% and 9.87  $\pm$  0.68% on the third and sixth day post WSSV infection, which were 1.94-fold (P < 0.05) and 2.69-fold (P < 0.01) higher than the Tris-HCl group in SGC of crayfish, respectively (Fig. 3). In mitochondrial permeability transition (MPT), the inner and outer mitochondrial membranes communicate via high-conductance channels that dissipate  $\Delta \psi$  and release cytochrome c. Induction of MPT can be caused by diverse stimuli and is believed to be involved in cell death [27]. In SGC stained with JC-1,



**Fig. 6.** The quantification of Bax inhibitor-1 transcript relative to 18s RNA transcript in the seven groups (Blank, Tris-HCl, WSSV, Tris-HCl-Tris-HCl, LY294002-Tris-HCl, Tris-HCl-WSSV and LY294002-WSSV) on the third, sixth and tenth day. Each symbol and vertical bars represented the mean  $\pm$  SD (N = 5). Significant differences between Tris-HCl and WSSV, Tris-HCl-Tris-HCl and LY294002-WSSV are indicated with one asterisk at P < 0.05 and two asterisks at P < 0.01.

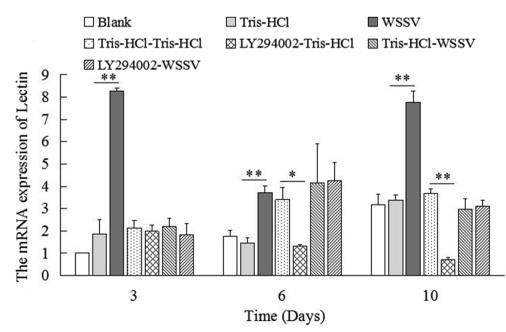


**Fig. 7.** The quantification of Toll transcript relative to 18s RNA transcript in the seven groups (Blank, Tris-HCl, WSSV, Tris-HCl-Tris-HCl, LY294002-Tris-HCl, Tris-HCl-WSSV and LY294002-WSSV) on the third, sixth and tenth day. on the third, sixth and tenth day. Each symbol and vertical bars represented the mean  $\pm$  SD (N = 5). Significant differences between Tris-HCl-Tris-HCl and LY294002-Tris-HCl were indicated with one asterisk at P < 0.05 and two asterisks at P < 0.01.

there was lower  $\Delta \psi$  (P < 0.01) in crayfish post WSSV infection compared to that in crayfish after Tris-HCl injection on the third day (Fig. 4). The result demonstrated that WSSV induced a stress response (SGC apoptosis), which might participate in the maintenance of the long-term WSSV infection and decline of the virus clearance. Targeting the PI3K/AKT pathway is a notable strategy taken by various viruses to inhibit apoptosis. For example, inhibition of PI3K activation with LY294002 could increase the cleavage and release of caspase-3, and induce apoptosis during largemouth Bass virus (LMBV), dengue virus (DEN) or Japanese encephalitis virus (JEV) infection [11-13]. However, our findings are in conflict with these results, which have suggested that PI3K inhibition is required to block apoptosis in WSSV-infected crayfish (Figs. 3 and 4). Similarly, in herpes simplex virus-1 (HSV-1)-infected OC3 cells, LY294002 treatment did not increase apoptosis and affect cell viability [28]. Therefore, the relationship between PI3K activity and apoptosis may vary in different host or virus strains. WSSV infection leads to the outbreak death of cultured Prawn in a short time, but crayfish is resistant and tolerance to WSSV

infection. Therefore, the balance between the pro- and anti-apoptosis activation processes regulated by PI3K/AKT signaling pathway would determine the fate of the WSSV-infected host.

Bax, a pro-apoptotic member of the Bcl-2 family, localizes in the cytoplasm but redistributes to mitochondria in response to apoptotic stimuli, where it induces cytochrome release [29]. The PI3K/AKT signaling pathway plays a pivotal role in the regulation of apoptosis, and AKT activates translocation of Bax from the cytoplasm to the mitochondria [30]. In order to investigate the mechanism of apoptosis in WSSV infected crayfish, Bax and Bax inhibitor-1 were examined. The expression level of Bax mRNA was increased on the third day after WSSV infection, which indicated that WSSV induced Bax production and Bax redistributed in the mitochondria, resulting in cytochrome release and apoptosis. A significant increase of Bax expression level in LY294002 pretreatment crayfish and a significantly lower proportion of apoptosis cells were observed (Figs. 3 and 4). LY294002 treatment increased Bax translocation to the mitochondria and be anti-apoptosis



**Fig. 8.** The quantification of lectin transcript relative to 18s RNA transcript in the seven groups (Blank, Tris-HCl, WSSV, Tris-HCl-Tris-HCl, LY294002-Tris-HCl, Tris-HCl-WSSV and LY294002-WSSV) on the third, sixth and tenth day. Each symbol and vertical bars represented the mean  $\pm$  SD (N = 5). Significant differences between Tris-HCl and WSSV, Tris-HCl-Tris-HCl and LY294002-Tris-HCl were indicated with one asterisk at P < 0.05 and two asterisks at P < 0.01.

activation processes. Crayfish Bax inhibitor-1 protein has been proven as an anti-apoptotic factor, which suppresses programmed cell death and is beneficial for WSSV replication [31]. The Bax inhibitor-1 levels in LY294002 pretreatment group were significantly lower than those in Tris-HCl pretreatment crayfish on the third day regardless whether the crayfish were infected with WSSV (Fig. 6). Bax inhibitor-1 protein neutralizes, balances Bax activity and plays a protective function as the main suppressor of Bax-induced cell death in environmental responses [32]. The results together demonstrated the PI3K/AKT signaling pathway played an important role in the regulation of host cell survival through regulation of Bax expression levels.

Toll-like receptors (TLRs), a major family of PRRs, play an important role in innate host defense as well as in initiation of adaptive immune responses. PI3K inhibitors (i.e., wortmannin and LY294.002) could modulate TLR-induced cytokine production and enhance antitumor immunity in humans. In addition, previous studies implicated PI3K and its substrate AKT in the regulation of TLR signaling. In the present study, the Toll mRNA expression was significantly down-regulated in LY294002 pretreatment crayfish on the third, sixth and tenth day (Fig. 7), which indicated that PI3K signaling played a major role in inducing the up-regulation of Toll expression and activating TLR signaling. Lectin proteins family is one of the important components in the immune defense system of invertebrate. The lectin mRNA expression was significantly increased in haemocytes of crayfish after WSSV infection on the third, sixth and tenth day, indicating lectin involved in the immune defense responses against WSSV in crayfish. There was a significant decrease lectin expression level in LY294002-Tris-HCl group compared to that in Tris-HCl-Tris-HCl group on the sixth and tenth day (Fig. 8). The lectin expression in LY294002 pretreatment crayfish was totally consistent with the previous Toll expression. The PI3K/AKT signaling is involved in the regulation of Toll and lectin signaling outcomes, which induced and regulated innate immune response in crayfish.

In conclusion, the PI3K/AKT signaling pathway play positive roles in WSSV replication. Blockage of the PI3K/AKT signaling pathway could decrease the WSSV copy numbers and host cellular apoptosis in WSSV-infected crayfish. Furthermore, the PI3K/AKT signaling pathway could regulate the apoptosis-related factors of Bax and Bax-inhibitor-1, as well as the level of innate immune defense protein (Toll and lectin).

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