



Short communication

Neuroendocrine immune-regulatory of a neuropeptide *ChGnRH* from the Hongkong oyster, *Crassostrea Hongkongensis*

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ABSTRACT

It is increasingly appreciated that neuroendocrine-immune interactions hold the key to understand the complex immune system. In this study, we explored the role of a reproductive regulation-related hormone, GnRH, in the regulation of immunity in Hong Kong oysters. We found that *vibrio* bacterial strains injection increased the expression of *ChGnRH*. Moreover, *ChGnRH* neuropeptide promotes the phagocytic ability and bacterial clearance effect of hemocytes which regarded to be the central immune organ. The content of cAMP after incubation with *ChGnRH* peptide was increased, which could be blocked by adenylyl cyclase inhibitor SQ 22,536. Furthermore, the stimulated effect of *ChGnRH* peptide on the phagocytosis and bacterial clearance was also blocked by SQ 22,536, H89 and enzastaurin, strongly demonstrating that cAMP dependent PKA and PKC signaling pathway was involved in *ChGnRH* mediated immune regulation. In conclusion, this study confirms the presence of neuroendocrine-immune regulatory system in marine invertebrates, which contributes to understand the complexity of oyster immune defense system.

1. Introduction

The mammalian endocrine and immune systems interact and form a network to maintain the homeostasis of the body, which is bidirectionally regulated by signal molecules such as hormones and cytokines [1,2]. The cross-talk involving the endocrine and immune systems is now largely established, which named neuroendocrine-immune regulatory system (NEI system). It was reported classical hormones such as prolactin (PRL), growth hormone (GH) and even glucocorticoids (GC) could be produced by immune cells [3,4]. In fish, growth hormone (GH) enhances many aspects of immune functions including non-specific defences; cytotoxic, phagocytic, haemolytic and lysozyme activities [5,6]. GH has been found to increase particle ingestion by fish leucocytes *in vitro*, indicating an activation of phagocytic activity [7,8]. *In vivo* and *in vitro* administration of GH enhanced superoxide anion production as a killing mechanism following phagocytosis [8–10]. GH also increased mRNA levels of superoxide dismutase, which catalyzes the dismutation of superoxide into oxygen and hydrogen peroxide [11]. All of these suggests an immune role of endocrine involving in the cell immunity.

The gonadotropin-releasing hormone (GnRH), also known as luteinizing hormone-releasing hormone (LHRH), is a decapeptide (pyroGlu-His-Trp-Ser-Tyr-Gly-Leu-Arg-Pro-Gly-NH₂) that plays an essential role

in the regulation of reproduction [12]. GnRH is responsible for the synthesis and secretion of follicle stimulating hormone (FSH) and luteinizing hormone (LH) [13], which effects the gonadal steroid hormone production and gametogenesis [14]. GnRH is also considered as a neurohormone, mainly produced in the preoptic area of the hypothalamus, where contains most of the GnRH-secreting neurons. The length of GnRH subtype is almost decapeptides that contains the highly conserved amino acid sequences in the N-terminal (Glp-His-Trp-Ser) and C-terminal (Pro-Gly-Gly-NH₂) regions, indicating that this molecular feature is crucial for receptor binding and activation [15]. In mammals, GnRH is found in organs outside of the hypothalamus and pituitary, such as the lymphocytes of the spleen and thymus [16]. With the deepening research on the interaction between the nervous system and the immune system, the role of GnRH in immune system has attracted more attention. Treatment with various concentrations of GnRH increased interleukin-2 receptor γ -chain mRNA in a dose-dependent manner, indicating that GnRH may be involved in lymphocyte activation [17]. Splenic and thymus lymphocytes have the capacity to produce immunoactive GnRH, further strengthening an association between the endocrine and immune systems [18–20].

More recently, new GnRH homologs have been identified in a variety of invertebrates [21–23]. In mollusks, GnRH mRNA is widely expressed in the central nervous system and accessory sex organs [24]

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or oocytes [22]. The GnRH in octopus has a high expression level in brain and acts as a multifunctional modulatory factor in memory processing, sensory, movement, autonomic functions and feeding [25,26]. In the Pacific oyster, two GnRH-related peptides (CgGnRH-a and CgGnRH-G) are characterized by mass spectrometry, which are highly conserved with other mollusks or vertebrates [27]. However, it is reviewed that the functions of the invertebrate GnRH-like peptides are not necessarily related to reproduction [28]. Considering the relationship between the endocrine and immune system [5,29,30], and the fact that vertebrates GnRH have been proved to involve in the process of immune regulation, we hypothesize that the neuroendocrine immunoregulatory of GnRH may be traced to lower invertebrates, such as mollusks. The Hongkong oyster *Crassostrea hongkongensis*, as an important global economic shellfish, mainly depends on innate immune system to defend invading microorganisms. Phagocytosis is one conserved cellular immune response for host to against microbes, or even apoptotic and necrotic cells [31,32]. It is well documented in oyster that defense against foreign bacterial is mediated by phagocytic cells which circulates in hemolymph. In this study, we discuss the role of neuropeptide ChGnRH in oyster immune system and its influence on the ability of oyster hemocytes to engulf the fluorescent bacterial, in order to discuss the exist of cross-talk between endocrine and immune system in invertebrate.

2. Materials and methods

2.1. Oyster collection and reagent

The Hongkong oysters, *Crassostrea hongkongensis* (two years old with an average 100 mm shell length) were obtained from Zhanjiang, Guangdong Province, China, and maintained at 22–25 °C in tanks with re-circulating seawater before experiments. Naive and pathogen-free oysters were chosen for the experiments, independently of their genetic background. The oysters were fed twice daily on *Tetraselmis suecica* and *Isochrysis galbana*. They were held for two weeks prior to experimentation. ChGnRH peptide were synthesized by GL biochem Ltd. Forskolin, SQ 22,536, H89 and enzastaurin were purchased from Sigma-Aldrich.

2.2. Tissue collection and bacterial challenge

To analyze the tissue distribution of ChGnRH, total RNA was extracted from gill, mantle, adductor muscle, digestive gland, gonads and hemocytes of three healthy *Crassostrea hongkongensis* individuals. To study the immune response of the ChGnRH, one hundred oysters were randomly divided into 2 groups and placed in 2 tanks: the bacteria challenge and control group. Oysters were challenged by injecting 100 µl bacteria *Vibrio alginolyticus* or *Vibrio parahaemolyticus* (1×10^8) bacteria suspended with Phosphate buffer saline (PBS) into adductor muscles. The control groups were injected with equal volume of PBS. Hemolymph was collected at scheduled intervals (3, 6, 12, 24, 48, 72 h after pathogenic challenge) from the pericardial cavity through the adductor muscle and immediately centrifuged (700 × g for 10 min at 4 °C) to separate the hemocytes from plasma. Four individuals were randomly sampled in each group at every time point after injection.

2.3. RNA extraction and cDNA synthesis

Total RNA was isolated from frozen tissue using TRIzol Reagent (Invitrogen, USA) according to the manufacturer's directions, and quantified by measuring absorbance at 260 nm. The integrity of RNA was checked by agarose gel electrophoresis. Purified RNA sample were diluted to 1 µg/µl and pooled to perform cDNA synthesis using PrimerScript™ First Strand cDNA Synthesis kit (TAKARA Bio Inc. Japan) following manufacturer's protocol.

2.4. Bioinformatics analysis of ChGnRH

Amino acid sequence of ChGnRH was aligned with GnRH protein sequences of different species obtained from BLAST analysis using BioEdit Sequence Alignment Editor. The phylogenetic tree was constructed using MEGA (Molecular Evolutionary Genetic Analysis) version 3.0 with the neighbor-joining method with 1000 bootstrap replicates.

2.5. Transcription analysis of ChGnRH by quantitative real-time PCR (qRT-PCR)

qRT-PCR analysis was used to determine the expression of ChGnRH mRNA in various tissues and during bacterial challenge using the gene-specific primers (Supp Table 1). Each assay was performed in triplicate with β-actin mRNA as internal control, as it is a gene which is least affected by bacterial infection and is independent of tissue type [33]. The qRT-PCR was conducted using a LightCycler 480 (Roche) in a reaction volume of 10 µl containing 1 µl of template cDNA, 5 µl of 2 × SYBR Green Mix, 1 µl of each primer (10 pmol/µl), and 3 µl of PCR-grade water. The qRT-PCR cycle program consist of one cycle of 95 °C for 1 min, following by 40 cycles of amplification 95 °C for 15 s, 55 °C for 15 s, 72 °C for 20 s, 85 °C for 20 s for signal collection in each cycles. The dissociation curve analysis of amplification products was performed to confirm the specificity at the end of each PCR reaction. The relative expression of ChGnRH gene was calculated using the $2^{-\Delta\Delta CT}$ method [34]. All data are given in terms of relative mRNA, expressed as means ± standard error of mean (SE).

2.6. Phagocytosis assay

Vibrio alginolyticus and *Vibrio parahaemolyticus* labeled with FITC (fluorescein isothiocyanate) could emit green fluorescence. Then the hemocytes were incubated with GnRH peptide for one hour. The bacteria *V. alginolyticus* and *V. parahaemolyticus* were incubated with the hemocytes cultured in the 24 well plates for 15 min in a 50:1 ratio. Next, hemocytes were washed three times with Tris buffer (pH8.0, 50 mM) and suspended in PBS + 15% EDTA solution. Finally, flow cytometry assay was performed to detect the phagocytosis of oyster hemocytes. Hemocytes phagocytosis was monitored using at least a total of 10,000 events.

Moreover, SQ 22,536 (500 µM), H89 (50 µM), enzastaurin (25 µM) were added to hemocytes with ChGnRH peptide (10^{-5} g), respectively and incubated for two hours, following by detecting the hemocytes phagocytosis as described above.

2.7. Bacteria clearance assay

The bacterial clearance assay was performed as described previously with some modification [35,36]. Two strains, *Vibrio parahaemolyticus*, *Vibrio alginolyticus*, (the concentration is probably OD = 0.2) were first prepared. Oyster hemocytes were collected and cultured in 24-well plates ($\sim 2.5 \times 10^5$ cells) for an hour with GnRH peptides, and then exposed to bacteria in a 50:1 ratio. After half hour incubation, extracellular bacteria were removed by washing with 0.02% trypsin-EDTA for four times. Immediately, the hemocytes were lysed in 1 mL PBS buffer containing 0.05% Triton X-100. Finally, 50 µl of lysate was used for the purpose of culturing on LB plates to achieve bacterial counts. For each group, three wells were used to perform bacteria clearance assay and each experiment was repeated three times. The survival rates of *Vibrio* was calculated by the formula: the survival rates = the number of survival bacterial/the total number of adding bacterial, which was reported previously [37].

Moreover, SQ 22,536 (500 µM), H89 (50 µM), enzastaurin (25 µM) were added to hemocytes with ChGnRH peptide (10^{-5} g), respectively and incubated for two hours, following by detecting the bacterial

clearance rate as described above.

2.8. ChGnRH stimulated the production of cAMP in hemocytes

Cyclic AMP (cyclic adenosine 3', 5'-monophospho, MW 351.2) is one of the most important intracellular mediators. In order to examine the effect of ChGnRH on intracellular cAMP production, a cAMP - Gs dynamic assay kit (Cisbio, 62AM4PEB) was used. Oyster hemocytes were collected and cultured for half an hour, followed by incubation with different concentrations (10^{-7} g, 10^{-6} g, 10^{-5} g) of ChGnRH peptide for 2 h. ChGnRH peptide were synthesized by GenScript Biotech Corp.

Next, hemocytes were lysed for detection of the cAMP concentration according to manufacturer's instructions. Different concentrations of SQ 22,536 and ChGnRH peptide (10^{-5} g) were added to hemocytes and incubated for two hours, following by detecting the cAMP concentration as described above.

3. Result

3.1. Sequence analysis and phylogenetic analysis of ChGnRH

Amino acid sequence alignments (Supp Fig.1A) of the GnRH precursor protein shows that 11 amino acids form the mature peptides and have a high degree of identity with the GnRH in mammals, amphibians, fish and other shellfish. Phylogenetic tree was generated based on the GnRH precursor protein from eleven species, including *Homo sapiens*, *Gallus gallus*, *Mus musculus*, *Sus scrofa*, *Sparus aurata*, *Xenopus laevis*, *Oryzias latipes*, *Danio rerio*, *Mizuhopecten yessoensis*, *Crassostrea virginica* and *Crassostrea hongkongensis* (Supp Fig.1B). The evolutionary tree revealed GnRH had one ancient origin prior to divergences between protostomia-deuterostomia.

3.2. Tissue distribution and temporal expression of ChGnRH following bacterial challenge

qRT-PCR was used to detect the distribution of ChGnRH mRNA in different tissues. The results revealed that ChGnRH was constitutively expressed in all tissues including mantle, gill, digestive gland, hemocytes, heart, adductor muscle and gonads, supporting that ChGnRH had a broad distribution range in cells and tissues [38,39]. Moreover, the expression level of ChGnRH was higher in hemocytes and gill than gonads as show in Supp Fig.2.

Next, we investigated the temporal mRNA expression of ChGnRH in the hemocytes post bacterial challenge (Fig. 1). After injection with *Vibrio alginolyticus* and *Vibrio parahaemolyticus*, the expression level of ChGnRH was up regulated within 24 h, and it was significantly changed at 6, 12 and 24 h. The highest expression level of ChGnRH could be observed at 12 h and 24 h respectively, post *V. alginolyticus* or *V.*

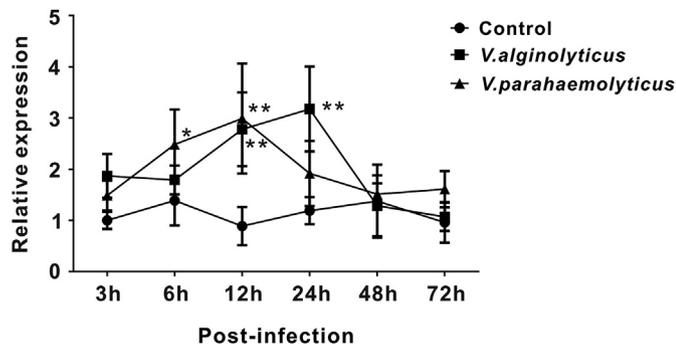


Fig. 1. ChGnRH mRNA expression after bacterial challenge in hemocytes. ChGnRH mRNA expression of each tissue was calculated by the $2^{-\Delta\Delta CT}$ method using β -actin as the internal control. Error bar represents the means \pm SE (N = 4).

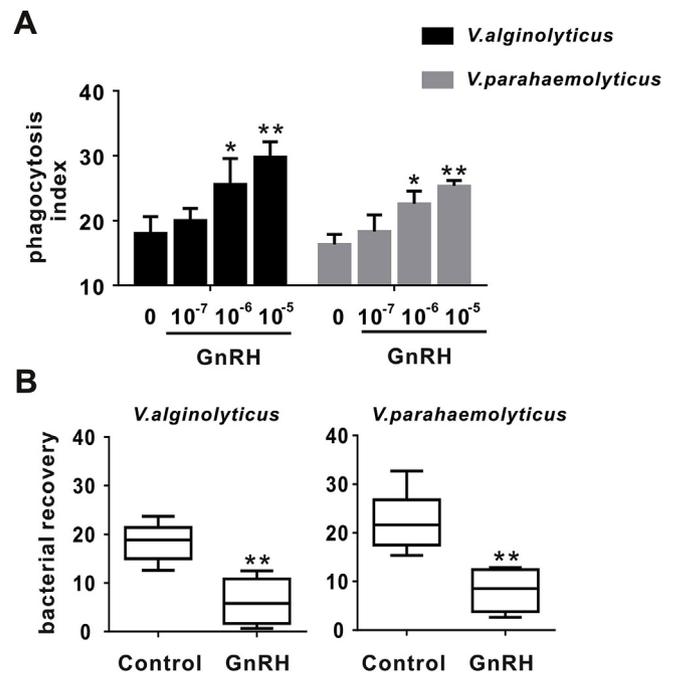


Fig. 2. ChGnRH peptide regulates innate immunity in hemocytes. (A) Data analysis was performed using GraphPad Prism 5 software and vertical bars represent the mean \pm SE (N = 3). The phagocytosis percentage of *v. alginolyticus* (left panel) and *v. parahaemolyticus* (right panel) of the ChGnRH (10^{-7} ~ 10^{-5} μ M) incubation group are higher than those of the control group. (B) The clearance capacity of hemocyte to bacteria (*Vibrio parahaemolyticus*, and *Vibrio alginolyticus*) at treatment of 10^{-6} μ M ChGnRH. Data represent the mean \pm SE (N = 3). Significant differences were indicated with * at $P < 0.05$ and ** at $P < 0.01$.

parahaemolyticus infection. Then the expression level recovered to the rest state. The control group injected with PBS maintained the normal expression.

3.3. ChGnRH promotes bacteria phagocytosis and clearance

Phagocytosis is a basic strategy of immune defense for oyster hemocytes. To investigate the possible effect of GnRH on the phagocytic ability of oyster hemocytes, the synthesized ChGnRH peptide was incubated into primary hemocyte culture. As shown in Fig. 2A, the results showed that the phagocytic ability was significantly increased to either *V. alginolyticus* or *V. parahaemolyticus* after incubation with ChGnRH dose of 10^{-6} and 10^{-5} μ M, indicating the key role of ChGnRH in the process of hemocytes immunity. Meanwhile, the dose-dependent effect of ChGnRH on the enhancement of hemocytes phagocytosis were also observed and the highest phagocytic proportion to *v. alginolyticus* and *v. parahaemolyticus* the reached 29.8% and 26.2% at dose of 10^{-5} μ M GnRH, which increased 1.7- and 1.6-fold when compared to control, respectively.

Bacterial clearance is an effective way of hemocytes to eliminate invaders pathogens. The results found that ChGnRH significantly enhanced the clearance of *v. alginolyticus* and *v. parahaemolyticus* (Fig. 2B). After 30min of hemocytes phagocytosis, the viable *v. alginolyticus* and *v. parahaemolyticus* occupies 18.4% and 21.8% of the bacteria amount initially added, which means that majority of bacteria were cleared by hemocytes. In the experimental group incubated with ChGnRH, only 6.2% and 8.5% of *v. alginolyticus* and *v. parahaemolyticus* survived.

3.4. ChGnRH enhanced the production of cAMP in the hemocytes

To investigate the signaling pathway of ChGnRH in oyster hemocytes, we examined its activation effect on cAMP production. Forskolin

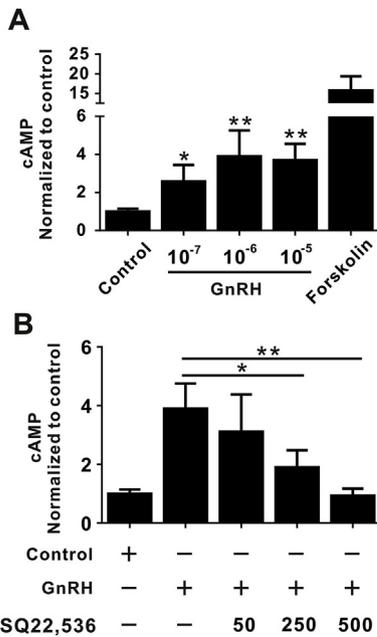


Fig. 3. *ChGnRH* peptide stimulates cAMP production in hemocytes. (A) The effect of *ChGnRH* peptide on the cAMP production at different concentration ($10^{-7} \sim 10^{-5}$ M). Treatment of forskolin (5 μ M) was as a positive control to stimulate the cAMP production. The adenylyl cyclase inhibitor SQ 22,536 (50–500 μ M) was used to block cAMP production. Data represent the mean \pm SE (N = 3). Significant differences were indicated with * at P < 0.05 and ** at P < 0.01.

is one adenylyl cyclase agonist to stimulate the cAMP production. After half of hour stimulation by *ChGnRH*, the hemocytes lysis was collected to detect the concentration of cAMP. As shown in Fig. 3A, it is clearly observed that the content of the cAMP was dose-dependent and significantly enhanced as the concentrations of *ChGnRH* increases, which reached 2.8-fold–3.9-fold higher than that of the control group. Meanwhile, the cell-permeable adenylyl cyclase inhibitor SQ 22,536 was co-incubated and significantly blocked the stimulation effects of *ChGnRH* in the dose dependent effects (Fig. 3B), confirmed cAMP production in GnRH stimulated hemocytes.

3.5. *ChGnRH* enhanced the hemocytes immunity via cAMP signaling pathway

In order to further investigate which cAMP signaling pathway involves *ChGnRH* mediated immune defense oysters, the selective inhibitor of cAMP-dependent protein kinase (PKA)-H89 and the protein kinase C beta inhibitor-enzastaurin, were used to co-incubation with *ChGnRH*, respectively. As shown in Fig. 4A, *ChGnRH* significantly enhanced the phagocytosis percentage of *v. alginolyticus* and *v. parahaemolyticus* and reached at 26.7% and 23.8%, and co-incubation with SQ 22,536 dramatically decreased the phagocytosis capability. Meanwhile, both H89 and enzastaurin could effectively block the stimulated effect of *ChGnRH* on phagocytosis, suggesting involvement of both PKA and PKC signaling pathway in immune defense. Next, we also test whether cAMP pathway regulates the ability of bacterial clearance in oyster. Similarly, *ChGnRH* could enhance the bacterial clearance to either *v. alginolyticus* or *v. parahaemolyticus*. However, this enhancement was blocked when co-treatment with SQ 22,536, H89 (50 μ M) and enzastaurin (25 μ M) (Fig. 4B). Taken together, these experiments demonstrated that cAMP dependent PKA and PKC signaling pathway was involved in *ChGnRH* mediated immune regulation.

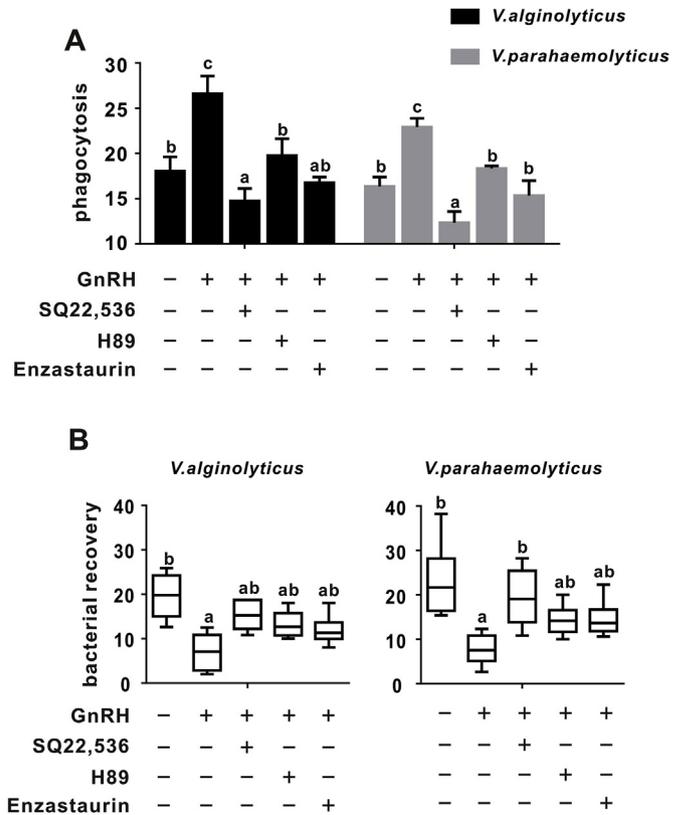


Fig. 4. cAMP dependent signaling pathway involves in *ChGnRH* mediated immune defense. Effects of SQ22,536, H89 and enzastaurin on *ChGnRH* stimulated phagocytosis (A) and bacterial clearance (B). Data represent the mean \pm SE (N = 3). Different letters refers to significant differences.

4. Discussion

Bidirectional communication between the immune and neuroendocrine systems is responsible for the infection with complex adaptations. Gonadotropin-releasing hormone (GnRH) is a highly conserved decapeptide hormone that plays a key role in the neuroendocrine system of vertebrates and also participates in the host's immune response process. It is known to possess direct immunomodulatory effects *in vivo* and *in vitro* in castrated rats and have been shown to exist in spleen and thymus, resembling a cytokine [18,40]. The presence of GnRH in human peripheral T-cells (CD4⁺, CD8⁺) has also been extensively reported, which suggest a potential role of GnRH in immune regulation [41]. Similar studies about cross-talk between endocrine and immune system have also been conducted in many marine invertebrate [42,43]. Our study reveals that *ChGnRH* has a high expression level in oyster immune organs, indicating a cross-talk between endocrine and immunity in mollusk.

Advances in molecular approaches in immunology prove that cross-talk involving the endocrine and immune systems is now largely established in vertebrate and invertebrate including oyster. However, the signaling pathway involved in GnRH dependent immune activation are still unclear. Generally, GnRH could stimulate cAMP second messenger pathway in varied cell types [44,45]. In agreement with this, our study was undertaken and demonstrated that low dose of Hongkong oyster GnRH peptide could elicit cAMP synthesis in the hemocytes cells, while adding SQ 22,536 (inhibitor of adenylyl cyclase) can effectively weaken the activation effect of *ChGnRH* on cAMP production. GnRH was also reported to increase cAMP levels and stimulated a cAMP-responsive promoter in rats that affected hormone release and synthesis [46]. The potential relationship of GnRH and cAMP has been strengthened recently observed, suggesting the conserved signaling pathway of GnRH

from human to oyster.

Cyclic AMP regulates a number of different cellular processes such as cell growth and differentiation, ion channel conductivity, synaptic release of neurotransmitters, and gene transcription [47]. Recent studies has demonstrated that activation of cAMP-dependent signaling pathways is necessary for control phagocytosis through actin rearrangements in human neutrophils [48], which plays a crucial role in engulfment of invaders and apoptotic cells, and maintaining the cell homeostasis [49,50]. GnRH was examined in phagocytic leucocytes of rainbow trout (*Oncorhynchus mykiss*), which indicated GnRH stimulates phagocytosis and superoxide production in fish leucocytes through a GnRH-receptor-dependent pathway [51]. Similar result is observed in Hongkong oyster that ChGnRH also harbors the capacity of improving phagocytosis and bacterial clearance in hemocytes to regulate the immune responses, which could be blocked through co-incubation of cAMP synthesis inhibitor SQ 22,536. So, these evidences strongly suggested that ChGnRH dependent phagocytosis activation is mediated through cAMP signaling pathway in oyster hemocytes. Downstream cAMP signaling is transduced by its interactions with effector molecules, such as protein kinase A (PKA), which increased expression of pro-inflammatory mediators [47,52]. cAMP analogues mediate the translocation of PKC to the nucleus and function in the regulation of gene expression [53]. Inhibition of PKA in Hongkong oyster with H89 has effect on the ability of bacterial clearance and phagocytosis. The same phenomenon could be observed when PKC was blocked by enzastaurin. These findings lead us to hypothesize that GnRH could constitute a mechanism for hemocytes phagocytosis and bacterial clearance via cAMP-dependent PKA/PKC signaling pathway.

In conclusion, this study for the first time confirms the presence of GnRH dependent immune regulatory system in the Hongkong oyster. GnRH could strongly promotes the hemocytes phagocytosis and bacterial clearance through cAMP and its downstream signaling pathways, PKA and PKC. Therefore, the cross-talk between endocrine and immune system in invertebrate not only benefits to revealing the complexity of neuroendocrine immunology in marine invertebrates, and contributes to new strategies in disease control in oyster aquaculture.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.fsi.2019.05.055>.

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