

Immune responses of white leg shrimp, Litopenaeus vannamei (Boone, 1931) to bacterially expressed dsRNA specific to VP28 gene of white spot syndrome virus

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32 Abstract

In the present study, dsRNA specific to VP28 gene of white spot syndrome virus 33 (WSSV) of shrimp was synthesized in Escherichia coli in large-scale and studied the immune 34 response of shrimp to dsRNA-VP28. The hematological parameters such as clotting time and 35 total haemocytes counts, and immunological parameters such as prophenoloxidase (proPO), 36 superoxide dismutase, superoxide anion and malondialdehyde content, as well as the mRNA 37 expression of ten immune-related genes were examined in order to estimate the effect of dsRNA-38 VP28 on the innate immunity of *Litopenaeus vannamei*. The activities of proPO, SOA and SOD 39 significantly increased in haemocyte after dsRNA-VP28 treatment, whereas MDA content did 40 not change significantly. Among the ten immune related genes examined, only the mRNA 41 expression of proPO, cMnSOD, hemocyanin, crustin, BGBP, lipopolysaccharides (LPs), lectin 42 and lysozyme in haemocyte, gill and hepatopancreas of L. vannamei, was significantly up-43 regulated at 12 h after dsRNA-VP28 treatment, while no significant expression changes were 44 observed in toll receptor and tumor receptor genes. The increase of proPO and SOD activities, 45 and SOA level and mRNA expression level of proPO, cMnSOD, hemocyanin, crustin, BGBP, 46 47 LPs, lectin and lysozyme after dsRNA-VP28 stimulation indicate that these immune related genes were involved in dsRNA-VP28-induced innate immunity in shrimp. 48

49 Keywords: *Litopenaeus vannamei*, Double-stranded RNA, Immunological parameters,
50 Hematological parameters, Immune genes expression.

51

52 **1. Introduction**

53 White spot syndrome virus (WSSV) is one of the most devastating shrimp viral pathogens and has a wide host range of several crustacean species, including shrimp, crab, and 54 55 crayfish. WSSV is responsible for 100% mortality within a few days after onset of the infection and is a serious threat to the shrimp culture industry worldwide (Lightner, 1996). Transmission 56 of the virus is mainly through oral ingestion and waterborne routes in farms and vertical 57 transmission in the case of shrimp hatcheries (Rosenberry, 2002). Considering the global 58 59 economic and sociological importance of shrimp farming, development of new control measures 60 becomes inevitable against the spread of WSSV. Protective effects of various immunostimulants such as oral administration of peptidoglycan (Itami et al., 1998), lipopolysaccharide (Takahashi 61 62 et al. 2000; Tsutsui et al. 2007), glucan (Song et al. 1997; Chang et al. 2003), and aqueous

extract of Cynodondactylon (Balasubramanian et al. 2007) have been reported against WSSV 63 infection. In contrast to the well studied effect of microbial immunostimulants on the immune 64 system of shrimp (Liu et al. 2005; Perez et al. 2005), Venegas et al. (2000) have also reported a 65 new concept of quasi-immune response against WSSV in *Penaeus japonicus*. Various vaccines 66 like inactivated WSSV vaccine (Namikoshi et al. 2004; Huang et al. 2005), antibacterial 67 components (George et al. 2006), and subunit recombinant vaccines (Kim et al. 2004; Li et al. 68 2005, 2006; Wei and Xu 2005; Jha et al. 2006) have also been tried so far against WSSV with 69 notable results. Because crustaceans have only a nonspecific innate immune response and no 70 long memory (Warren et al. 2006), developing a vaccine and using it as an effective control 71 strategy is difficult. 72

Double-stranded RNA (dsRNA) is an intermediate produced in the replication of many 73 viruses, which induces multifaceted immune responses in mammals, including RNA interference 74 (RNAi) and the production of interferon (IFN) (Kapadia et al., 2003). RNAi is a sequence 75 dependent antiviral mechanism, which can inhibit the replication of virus by suppressing the 76 expression of virus genes in host cells (Li et al., 2002; Silhavy et al., 2002; Li et al., 2004). In 77 78 addition to RNAi, another sequence-independent immune mechanism, IFN-related immune responses are also induced by dsRNA in mammals (Robalino et al., 2004). In this process, 79 80 dsRNA is recognized by Toll like receptor 3 (TRL3) which then activates innate antiviral immune responses via My88-dependent and independent pathways, and finally leads to the 81 82 production of interferons (IFNs) (Robalino et al., 2004). IFNs induce the expression of many proteins including RNA-dependent protein kinase (PKR), the Mx (myxovirus) protein and 83 84 oligoadenylatesynthetase, which have been demonstrated to play important roles in antiviral immunity. Moreover, dsRNA can also directly activate intracellular PKR and induce antiviral 85 86 responses. Activated PKR inhibits cellular and viral protein synthesis via phosphorylation of eukaryotic translation initiation factor 2a (eIF2a) which further inhibits the replication of virus 87 (Robalino et al., 2004; Meurs et al., 1990). 88

The RNAi pathway is thought to be an ancient mechanism for protecting the host and its genome against viruses and rogue genetic elements by the process of mRNA degradation that is induced by double-stranded RNA (dsRNA) in a sequence-specific manner (Jones et al. 1999; Zamore et al. 2000; Bernstein et al. 2001). Several recent papers have reported that siRNAs (Westenberg et al. 2005; Xu et al. 2007) or dsRNA synthesized via *in vitro* methods (Robalino et

al. 2004; Robalino et al. 2005; Kim et al. 2007) serve as potential therapeutic agents for treating 94 white spot syndrome disease. In the marine shrimp *Litopenaeus vannamei*, the antiviral response 95 can be induced by sequence-independent or sequence-specific dsRNA, which may activate 96 RNAi-like mechanisms (Robalino et al. 2004, 2005). Westenberg et al. (2005) have used siRNAs 97 specific to the VP15 and VP28 genes of WSSV and observed that shrimp injected with VP15 or 98 VP28 siRNASs before WSSV challenge had a significantly lower mortality. Xu et al. (2007) 99 100 have used a specific 21-bp short interfering RNA targeting the VP28 gene of WSSV and the results revealed that the transcription and expression of the VP28 gene were silenced. The 101 introduction of long dsRNAs corresponding to viral proteins is very effective in blocking WSSV 102 infection in *Penaeus chinensis* (Kim et al. 2007). Yodmuang et al. (2006) reported the protection 103 of P. monodon against yellow head virus (YHV) infection by silencing the genes of YHV using 104 105 long dsRNA encoding structural and non-structural proteins of YHV. Recently, Sarathi et al. (2008, 2008a) reported the possibility of protecting Penaeus monodon against white spot 106 syndrome virus (WSSV) infection via RNAi technology by oral administration of bacterially 107 expressed VP28 (VP28dsRNA) gene of WSSV. 108

109 The detailed mechanism of this dsRNA-induced antiviral immunity is still not well understood. In present study, bacterially synthesized dsRNA specific to VP28 gene of WSSV 110 111 promoted gene-specific interference with the WSSV infection in shrimp and the effect of dsRNA-VP28 on the immune response of L. vannamei was explored for obtaining more insight 112 113 into the mechanism of dsRNA-VP28 induced immunity in shrimp. The temporal activity changes of superoxide dismutase (SOD), prophenoloxidase (proPO), superoxide anion (SOA) and 114 115 malondialdehyde (MDA) variation; hematological parameters such as clotting time and total haemocytes count (THC) changes, as well as the mRNA expression of some immune-related 116 117 genes in L. vannamei were examined after dsRNA-VP28 treatment to estimate the effect of dsRNA-VP28 on the immune system to determine the dsRNA-VP28 induced immune responses 118 in L. vannamei. 119

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121 **2.** Materials and Methods

122 **2.1.** Collection of experimental animals

123 White leg shrimp, *L. vannamei* (10–15 g body weight), were collected from 124 Nagapatinam, Tamilnadu, Indiaand were maintained in 1000-l fiberglass tanks with air-lift

biological filters at room temperature (27–30 °C) with salinity between 20 and 25 parts per
thousand (ppt) for 1 week prior to experiment. During the experiment, the shrimp were fed with
commercial pellet feed (CP feed, Thailand), and water was totally exchanged daily.

128 2.2. Isolation of dsRNA-VP28

dsRNA-VP28 was synthesized according to the method of Sarathi et al. (2008) with 129 slight modification. For the isolation ofdsRNA-VP28, IPTG-induced bacterial cells were pelleted 130 for 15min at 3000 g (4 °C) and resuspended in 1.25 ml of TE buffer (10 mM Tris-HCl, 1 mM 131 EDTA, pH 7.5). A minimum volume of 0.75 ml of the cell suspension was replenished with an 132 equal volume of phenol-chloroform-isoamyl alcohol and heated at 65°C for 10 min. The 133 samples were then centrifuged for 10 min at 13,000 rpm and the aqueous phase transferred into a 134 fresh tube. Nucleic acid was then precipitated with isopropanol and kept at -20° C for 10 min and 135 then centrifuged. Supernatant was discarded and the pellet was washed with 70% ethanol. After 136 drying, the pellet was resuspended in 100 µl of Tris–NaCl–MgCl₂–dithiothreitol buffer. Eighteen 137 microliters of this solution was supplemented with 1 µl of DNaseI (3 mg/ml) and digested for 15 138 min at room temperature to remove DNA, followed by extraction with phenol/chloroform and 139 precipitation with ethanol. The pellet was resuspended in 50 µL RNase-free PBS buffer and 140 subjected to electrophoresis for RNA integrity confirmation. The dsRNA concentration was 141 quantified by measuring the absorbance using UV spectrophotometer at the wavelength of 260 142 and adjusted to a final concentration of 1 µg/ml. dsRNA was injected intramuscularly at the third 143 144 abdominal segment of shrimp using 1 mL syringe.

145 2.3. Injection of dsRNA-VP28 in shrimp

146 The shrimp were divided into two groups and maintained in aquarium tanks (10shrimp/tank) as described above. In the Group I, the shrimp were injected intramuscularly 147 148 with 50 μ L dsRNA at the concentration of 1 μ g/ μ l specific to VP28 gene of WSSV at third abdominal segment of shrimp. In Group II, the shrimp were injected intramuscularly with 50 µL 149 150 PBS (pH 7.0) and this group of shrimp was used as negative control. During the experimental period, three shrimp were randomly collected from each group at 0 h, 3 h, 6 h, 12 h, 24 h and 48 151 152 h post-injection. The haemolymph was drawn from the heart using a 23 gauge needle attached to a 1 mL syringe. After collection, the haemolymph was quickly mixed with ice cold anticoagulant 153 solution (27 mM Na citrate, 336 mM NaCl, 115 mM glucose, 9 mM EDTA, pH 4.6 at 28 °C). 154 Samples were immediately centrifuged at 800 x g at 4°C for 10 min to collect the haemocytes. 155

156 Cell pellet was gently washed with 500 μ L of wash solution (415 mmol/L NaCl, 100 mmol/L 157 glucose, pH 7.0) and centrifuged again at 800 g for 5 min. The haemocytes were resuspended in 158 1 mL lysis buffer (415 mmol/L NaCl, 100 mmol/L glucose, 10 mmol/L cacodylic acid, 5 159 mmol/L CaCl₂, pH 7.0) and sonicated at 40% power for 2 min. Cell debris was removed by 160 centrifuging at 13,000 g for 10 min, and the resultant haemocyte lysate supernatant was used for 161 analyzing proPO, SOD and SOA activity and MDA content.

162 **2.4. Immunological parameters**

163 **2.4.1. Prophenoloxidase activity**

Prophenoloxidase activity of PBS or dsRNA-VP28 injected shrimp was analyzed 164 spectrophotometrically by the method of Hernandez-Lopez et al. (1996) by recording the 165 formation of dopachrome produced from L-dihydroxyl phenylalanine (L-DOPA). Briefly, 166 hemolymph collected from normal or experimental shrimp was diluted and centrifuged at 800 x 167 g at 4 °C for 20 min and the pellet was collected. Then the pellet was suspended gently in 168 cacodylate buffer (0.01 M sodium cacodylate, 0.45 M sodium chloride, 0.10 M trisodium citrate, 169 pH 7.0). The sample was again centrifuged as described above. After removing the supernatant, 170 the pellet was resuspended in 100 µL cacodylate buffer. The sample was incubated after adding 171 50 µL trypsin (Sigma, 1 mg ml⁻¹) at 25°C for 10 min, which served as an activator; 50 µL DOPA 172 was then added followed by 800 µL of cacodylate buffer 5 min later. The optical density was 173 measured at 490 nm wave length by UV–VIS spectrophotometer (Shimadzu, UV 2450). 174

175 **2.4.2. Superoxide anion detection**

Superoxide anion was quantified in PBS or dsRNA-VP28 injected shrimp by the method 176 177 of Song and Hsieh, (1994). Briefly, 100 µL of hemolymph was collected from PBS or dsRNA-VP28 injected shrimp and centrifuged at 800 g for 5 min. After centrifugation, the supernatant 178 179 was removed and 100 µL of HBSS was added to haemocytes including normal sample tubes and allowed to react for 30 min at 37°C. After incubation, haemocytes were washed three times with 180 181 HBSS and then stained with Nitro blue tetrazolium (NBT) solution $(0.3\%, 100 \,\mu\text{L})$ for 30 min at 37°C. Removing the NBT solution and adding absolute methanol stopped the staining reaction. 182 183 The stained hemocytes were washed three times with 70% methanol and air dried. Added 120 µL of 2 M KOH and 140 µL DMSO to the sample to dissolve the cytoplasmic formazan. The optical 184 density of the dissolved formazan was read on a UV–VIS spectrophotometer (Shimadzu, UV 185 186 2450) at the wave length of 630 nm.

187 **2.5.** Antioxidant parameters

188 **2.5.1.** Assay of superoxide dismutase

SOD activity was determined in PBS or dsRNA-VP28 injected shrimp by the method of 189 190 Beauchamp and Fridovich (1971) using NBT in the presence of riboflavin. Briefly, 100 μ L of hemolymph was obtained from PBS or dsRNA-VP28 injected shrimp and homogenized in a 191 mechanical homogenizer containing 0.5 mL of phosphate buffer (50 mM, pH 7.8). The 192 193 homogenate was centrifuged for 5 min at 5700 g at 4°C and the supernatant recovered was heated for 5 min at 65°C to obtain a new supernatant after centrifugation, which was stored at 194 -20 °C until use. Samples were maintained on ice at all times to avoid protein denaturation. A 195 mixture of NBT, 20 µM of reaction mixture (0.1 Mm EDTA, 13 µM methionine, 0.75 mM NBT 196 and 20 µM riboflavin in phosphate buffer 50 mM, pH 7.8) and 100 µL of the crude extract was 197 placed under fluorescent light for 2 min or until A560 in normal sample tubes reached 0.20-0.25 198 OD. The results were expressed as relative enzyme activity. 199

200 2.5.2. Malondialdehyde content

In this method described by Beuge and Aust (1978), a mixture of 100 μ L Tris buffer (150mM, pH 7.1), 10 μ L ferrous sulfate (100mM), 10 μ L ascorbic acid (150 mM), 780 μ L distilled water and 100 μ L of cell extract (haemocytes) was incubated at 37°C for 15min. Thiobarbituric acid (0.375%, 2mL) was then added to the mixture and allowed to react at 100°C (in water bath) for 15min. The reaction mixture was then centrifuged (800×*g* for 10min) and supernatant was read at 532nm against blank. The MDA content was expressed as nmol per mg of protein (nmol/mg protein).

208 **2.6. Hematological assays**

209 **2.6.1.** Clotting time

210 The clotting time of hemolymph was measured according to the method described by Sachdev (1983). The hemolymph was collected from PBS or dsRNA-VP28injected shrimp 211 directly into a pre-cooled glass capillary tube of 1.2 mm diameter and 50 mm length. After the 212 hemolymph was inserted, the tube was kept at vertical position with the sample at the upper end. 213 214 Then the tube was maintained vertically until the gravity forced hemolymph to reach the lower end of the tube, after which the tube was turned to 180°. This was repeated until the hemolymph 215 clotted. The clotting time is defined as the point when the flow of hemolymph ceases and 216 217 presumably total clotting occurs. The clotting time was recorded.

218 **2.6.2. Total haemocytes count**

Total haemocyte count (cells/ml) was performed using a Burker hemocytometer as described by Sachdev (1983). One mL of hemolymph was drawn directly from the heart of shrimp injected with dsRNA-VP28 or normal shrimp by inserting a 23 gauge needle attached to a 2 ml syringe containing 1 ml of ice-cold anticoagulant Alsever solution (27 mM sodium citrate, 336 mM NaCl, 115 mM glucose, 9 mM EDTA, pH 4.6 at 28 °C). A drop of hemolymph with anticoagulant mixture was placed on a hemocytometer to measure the total haemocytes using a phase contrast microscope and counted manually in all 25 squares (0.1 mm³).

226 **2.6.3.** Oxyheamocyanin levels

For measurement of oxyheamocyanin level, 50 μ L of hemolymph from normal or experimental animal was immediately diluted with 900 μ l of distilled water in a 10-mm quartz cuvette and the absorbance was measured at 335 nm using a UV–VIS spectrophotometer (Shimadzu, UV 2450). The concentration of oxyhemocyanin was determined based on the method of Nickerson and Van Holde (1971) and Hagerman (1983).

232 2.7. Quantification of immune-related gene expressionby real-time PCR

233 The haemolymph, hepatopancreas and gill tissues were collected separately from each shrimp for extracting total RNA using Trizolreagent). Two mg of DNase-treated (RQ1 DNase I, 234 235 Promega) total RNA and oligo (dT)-adaptor primer were used to synthesize first-strand cDNA using M-MLV reverse transcriptase (NEB, USA). The cDNA was quantified using the 236 NanoDrop 2000C spectrophotometer (Thermo Scientific, Waltham, USA) at 260 nm. The 237 mRNA level of immune-related genes was determined using SYBR green quantitative real-time 238 239 PCR in ABI PRISM 7300 Sequence Detection System (ABI, USA). The amplification was performed in a 10 µL reaction volume containing 5 µL of 2 x SYBR Green Master Mix (ABI, 240 241 USA), 1µL of diluted cDNA, 0.5µL of each primer and 3.5 µL of DEPC-treated water. DEPCwater for the replacement of cDNA template was used as the negative control. The thermal 242 profile for the SYBR green real-time PCR was 50 °C for 2 min and 95 °C for 10 min followed 243 by 40 cycles of 95 °C for 15 s and 72 °C for 1 min. The expression of immune-related genes was 244 245 normalized to the expression of β -actin gene for each sample. The details of primersets used in 246 this assay are shown in Table 1. Data analysis was executed using mathematical model for relative quantification in real-time PCR by Pfaffl. (2001). ΔC_T represented the difference 247 between C_T for target gene and the internal control, and $\Delta\Delta C_T$ was obtained by subtracting the 248

249 ΔC_T for the control (PBS) from the ΔC_T for each sample. The obtained data were subjected to 250 statistical analysis.

251 **2.8. Data analysis**

Statistical analysis was executed using Excel software. All data obtained from the experiments were analyzed using one-way ANOVA (P<0.05 as significant level). Statistical calculations were performed using SPSS (version 16) software. Results are expressed as mean±SD. A significance level of p<0.05 was used for statistical testing.

256

257 **3. Results**

The gene encoding VP28 protein of WSSV was amplified by PCR and amplified product 258 was cloned into LITMUS38i vector. The recombinant plasmid bearing the VP28 gene was 259 transformed to E. coli HT115DE3 and dsRNA specific to VP28 was expressed under induction 260 of 1 mM IPTG. After induction with IPTG at 37°C, dsRNA was isolated from induced 261 LITMUS38i-VP28 E. coli HT115DE3. The isolated dsRNA was analyzed via agarose gel 262 electrophoresis (Fig.1). A band of dsRNA-VP28 corresponding to the molecular weight of 263 264 615 bp was observed in 1.2% agarose gel electrophoresis in the sample of IPTG-induced LITMUS38i-VP28-E. coli HT115DE3 (Fig.1, lane 3), whereas no dsRNA band was found at 265 266 the same position in both the non-induced LITMUS38i-VP28- E. coli HT115DE3 and induced LITMUS38i-E. coli HT115DE3 (Fig.1, lanes 1 and 2). The result of agarose gel indicates the 267 268 successful expression of dsRNA-VP28in E. coli.

In the experiment, no mortality was recorded in shrimp injected with dsRNA-VP28 and 269 270 significant immune response was observed when compared to shrimp injected with PBS. Significant change in the level of proPO, SOD and SOA was observed in shrimp injected with 271 272 dsRNA-VP28 when compared to shrimp injected with PBS (Figs.2, 3 and 4). The level of proPO activity increased to 8.3 Unit/mg protein in shrimp injected with dsRNA-VP28 at 24 h.p.i. 273 274 whereas the level was about 4.9 Unit/mg protein in shrimp injected with PBS (Fig. 2). The proPO activity gradually increased, reached highest level at 24 h p.i. in dsRNA-VP28 injected 275 276 shrimp. Significant change in the level of SOA and SOD was observed in shrimp injected with dsRNA-VP28 and the level was found to be the highest at 24 h p.i. when compared to the shrimp 277 injected with PBS (Figs. 3 and 4). The SOA level remained at a low level during the initial 6 278 hours and increased 3 to 4 folds after 12 h post injection of dsRNA-VP28. No significant change 279

280 was observed in MDA content in shrimp injected with dsRNA-VP28 in comparison with PBS-

injected shrimp (Fig. 5).

282 The clotting time of hemolymph was estimated in shrimp injected with dsRNA-VP28 and shrimp injected with PBS, and the results are shown in Fig. 6. There is no significant change in 283 the clotting time between shrimp injected with PBS and shrimp injected with dsRNA-VP28 at 284 different time interval after injection. Like clotting time, no significant change was observed in 285 total haemocyte count in shrimp injected with dsRNA-VP28 and normal shrimp (Fig.7). The 286 level of oxyhemocyanin (OHC) was determined in shrimp injected with dsRNA-VP28 and 287 shrimp injected with PBS at different time intervals. The results revealed that significant change 288 was found in shrimp injected with dsRNA-VP28 after 12 h p.i. The OD value increased to the 289 highest value of 0.61 in dsRNA-VP28 injected shrimp whereas the OD value for OHC in PBS 290 injected shrimp was found to be 0.342 at 24 h p.i. (Fig.8). 291

The mRNA expression level of ten immune-related genes was quantified in haemocytes, 292 hepatopancreas and gill tissue of shrimp injected with dsRNA-VP28 by quantitative real-time 293 PCR in order to investigate immune response todsRNA-VP28ds, and the results are presented in 294 295 Table 2. Lowered C_T value and enhanced mRNA expression level were observed in the genes of proPO, cMnSOD, hemocyanin, crustin, BGBP, lipopolysaccharides (LPs), lectin, lysozyme, toll 296 receptor and tumor receptor in haemocytes, hepatopancreas and gill tissue of dsRNA-VP28 297 injected L. vannamei (Figs. 9A - J). The β -actin gene served as internal control in the 298 299 haemocytes, hepatopancreas and gill of L. vannamei during the experiment. Among these immune-related genes, the expression level of proPO gene was enhanced about 12.17 fold, 8.9 300 301 fold and 5.8 fold in haemocytes, gill and hepatopancreas, respectively in dsRNA-VP28 injected shrimp (Fig.9A). In the case of cMnSOD gene, the expression level was increased about 10.2 302 303 fold, 4.8 fold and 5.4 fold in haemocytes, gill and hepatopancreas, respectively in response to dsRNA-VP28 injection in shrimp (Fig.9B). The expression level of hemocyanin gene was 304 enhanced about 6.01 fold, 5.1 fold and 3.9 fold in haemocytes, gill and hepatopancreas, 305 respectively in dsRNA-VP28 injected shrimp (Fig.9C). In the case of crustin gene, the 306 307 expression level was increased about 6.3 fold, 6.1 fold and 2.7 fold in haemocytes, gill and hepatopancreas, respectively in response to dsRNA-VP28 injection in shrimp (Fig.9D). The 308 enhanced mRNA expression level was observed in all the tissues tested in shrimp in response to 309 310 dsRNA-VP28 in the case of BGBP (Fig.9E), LPs (Fig.9F), lectin (Fig.9G) and lysozyme

311 (Fig.9H). But no change in the mRNA expression level was observed in the case of toll receptor

gene (Fig.9 I) and tumor receptor gene (Fig.9J) in response to dsRNA-VP28 injection in shrimp.

At 12 h, there was a significant (*P<0.05, dsRNA-VP28 treatment) increase in the relative abundance of proPO, cMnSOD, Hemocyanin, Crustin, BGBP, LPs, Lectin and Lysozyme genes mRNA. As time progressed, the expression of immune genes dropped gradually, and decreased to its original level at 48 h.post dsRNA-VP28 injection.

317

318 4. Discussion

Application of RNAi technology against WSSV is a possible solution to control the quick 319 spread of this deadly disease (Robalino et al., 2004, 2005; Westenberg et al., 2005; Xu et al., 320 2007; Kim et al., 2007). It is not possible to apply in vitro synthesized dsRNA and siRNA 321 available for RNAi therapy in the shrimp culture ponds, because they cannot be produced in 322 large quantities. dsRNA can stimulate different immune responses in mammals based on their 323 length (Dong et al., 2009). dsRNA shorter than 30 bp, called small interference RNA (siRNA), 324 induces RNA interference (Elbashir et al., 2001, 2001a) while dsRNA longer than 30 bp induces 325 326 the production of IFNs and therefore triggers strong antiviral immunity (Fire et al., 1998; Tuschl et al., 1999). In invertebrates, there is no dsRNA-induced IFN-related immunity because of the 327 328 lack of IFN-related genes (Robalino et al., 2004), and long dsRNA is often used to induce RNA interference (Williams et al., 1979; Misquitta and Paterson, 1999; Caplen et al., 2000; Robalino 329 330 et al., 2005). Recently, antiviral immune responses were observed in some crustacea after challenging with dsRNA (Robalino et al., 2004, 2005; Dong et al., 2009). But this antiviral 331 332 immunity was not as strong as it is in mammals and could be overcome by a high dose of infectious virus (Robalino et al., 2004). The mechanism of this immunity is suspected to be 333 334 different from the mechanism in mammals because there is no report of the presence of IFNrelated genes in invertebrates. Moreover, the antiviral immunity induced by dsRNA in 335 crustaceans is sequence-independent which is different from RNAi. 336

Changes in the level of prophenoloxidase activity observed in the present study indicate the involvement of dsRNA-VP28-induced immunity in shrimp as observed in crab by Dong et al. (2009). A significant elevation of proPO activity observed in the haemolymph of *L. vannamei* injected with dsRNA-VP28 might be due to the dsRNA treatment which triggered the prophenoloxidase activating system and transformed the prophenoloxidase into phenoloxidase.

In crustacea, melanization occurs when the cellular defense reactions are initiated (Ratcliffe et 342 al., 1985; Soderhall et al., 1986). Phenoloxidase, the key enzyme in the synthesis of melanin, 343 occurs in hemolymph as an inactive proenzyme prophenoloxidase (proPO) (Chang et al., 2003; 344 Balasubramanian et al., 2008). ProPO is activated to form phenoloxidase when it reacts with 345 zymosan (carbohydrates from yeast cell walls), bacterial lipopolysaccharide (LPs), urea, calcium 346 ions, trypsin, or heat (Soderhall et al., 1984, 1986). Results from several experiments have 347 confirmed that apart from their role in melanization, components of the putative proPO 348 activating system stimulate several cellular defense reactions, including phagocytosis, nodule 349 formation, encapsulation, and haemocytes locomotion (Soderhall et al., 1986; Johanson et al., 350 2000). The increase of phenoloxidase activity might defend the animal against invading 351 pathogens by promoting the melanization reaction, and generating highly cytotoxic quinones that 352 could inactivate viral pathogens or by enhancing other cellular activities such as phagocytosis 353 and encapsulation (Dong et al., 2009). This mechanism is especially important to protect the 354 shrimp against viral infection, because viral infection can inhibit the activity of phenoloxidase 355 activity in crustaceans (Mathew et al., 2007), while any reductions in the activity of 356 357 phenoloxidase activity might lead to the failure of phagocytosis (Mathew et al., 2007) and make the animals susceptible to infection. The dsRNA-induced mechanism help to improve the 358 condition of enhancement phenoloxidase activity and similar observation was reported in 359 Chinese mitten crab Eriocheirsinensisby injection of two double-stranded RNA (GFP-dsRNA 360 361 and NoPSD-dsRNA) (Dong et al., 2009). In addition, there are other reports about the enhancement of shrimp resistance to viral infection after stimulation with LPs (Takahashi et al., 362 363 2000), or 1, 3-glucans (Chang et al., 2003). The exact mechanism of dsRNA activation on the prophenoloxidase system still is not known and requires further investigation. 364

365 The superoxide anion activity increased significantly in dsRNA-VP28 administered shrimp when compared to normal control shrimp. The production of O₂⁻ has been reported as an 366 accurate method to measure the effectiveness of potential immunostimulants (Munoz et al., 367 2000). The higher level of O₂⁻ in dsRNA-VP28 administered shrimp indicates that dsRNA-VP28 368 369 may be the potential immunostimulant. Downs et al. (2001) reported that increase in the superoxide anion against pathogens is considered to be beneficial after exposing shrimp to 370 immunostimulants. These facts suggested that increase in the superoxide anion is considered to 371 372 be beneficial with respect to increased immunity.

373 Antioxidation systems play an important role in clearing excess ROS and protecting organisms from injury by ROS (Dong et al., 2009). In the present study, a significant increase of 374 375 SOD activity was observed in L. vannamei after dsRNA-VP28 injection, which indicated that this antioxidation system was involved in the dsRNA-VP28-induced immune response. The high 376 level SOD could convert superoxide anion into hydrogen peroxide, maintain the balance of 377 pro/antioxidant and protect cells against injury from lipid peroxidation. It is well known that 378 pathogen infection always results in inhibition of the SOD activity and a dramatic increase in the 379 level of ROS (Schwarz, 1996; Dong et al., 2009). High level ROS can lead to lipid peroxidation 380 of cell membranes or reduction in the PO activity (Mathew et al., 2007), and can result in high 381 mortality. The foreign stimuli enhancing the SOD activity will augment the host resistance to a 382 pathogen. The enhanced SOD activity in response to dsRNA-VP28injection indicates that 383 dsRNA-VP28 could strengthen the innate immunity in L. vannamei via this antioxidation system. 384

Malondialdehyde (MDA) content is an indicator of lipid peroxidation and is often used to 385 evaluate the damage of membranes resulting from oxidative stress. In the present study, MDA 386 387 content was also measured to examine the lipid peroxidation resulting from dsRNA-VP28 388 stimulus. There was no significant variation of MDA content in haemolymph of L. vannamei treated with dsRNA-VP28 compared to control animals as reported by Dong et el. (2009) in crab 389 injected with dsRNA. This suggests that dsRNA-VP28, although a viral-related molecular 390 patternhas no role in excessive ROS production as in the case of viral infection. This is 391 392 consistent with the previous report (Robalino et al., 2004) in which different doses of dsRNA did not result injury to shrimp. It is also supported by an earlier report that structural proteins of the 393 394 virus rather than dsRNA were responsible for ROS production in mouse splenocytes (Schwarz, 1996). 395

Balasubramaiyam et al. (2008) reported that there were no significant changes in clotting time (CT) and total haemocytes count (THC) between normal and plant (*Cynodondactylon*) treated shrimp at different day of post injection. Similar resultswere also observed in clotting time (CT), total haemocytes count (THC) and oxyhemocyanin of hemolymph in the present study. No significant changes were observed in clotting time, total haemocytes count and oxyhemocyanin level between control shrimp and dsRNA-VP28 treated shrimp.

Hepatopancreas and haemocytes which have a vital role in the immune response, are the major sites for the synthesis of immune defence molecules and are involved in eliminating 404 pathogens or other particulate matter (Johnson, 1987; Vogt, 1996; Gross et al., 2001). Gill is an important organ of respiration as well as of osmoregulation (Mantel and Farmer, 1983; Pequeux, 405 406 1995). These organs were selected for quantification of expression of immune genes in response to dsRNA-VP28 injection. One of the interesting hypotheses derived from our observation of 407 activation of RNAi and innate immunity by dsRNA-VP28 is that these two pathways interact 408 functionally to confer immunity to a viral pathogen. Such a possibility was previously 409 410 unrecognized among invertebrates, as L. vannamei is the only invertebrate in which dsRNA has been shown to induce both innate immune reactions and RNAi-like antiviral immunity (Robalino 411 et al., 2005). 412

Among the ten immune-related genes examined, proPO, cMnSOD, hemocyanin, crustin, 413 BGBP, LPs, lectin and lysozyme genes responded to dsRNA-VP28 challenge at different time 414 intervals in hemocyte, gill and hepatopancrease of shrimp by qPCR analysis. The expression of 415 these eight genes (proPO, cMnSOD, hemocyanin, crustin, BGBP, LPs, lectin and lysozyme) was 416 up-regulated by dsRNA-VP28 challenge, which indicated that these eight immune genes were 417 involved in the immune response of the *L. vannamei* in response to dsRNA-VP28. Another two 418 419 genes namely toll receptor and tumor receptor genes were not significantly up-regulated in dsRNA-VP28 treated shrimps. Recently, a TLR (Toll Receptor) was characterized in L. 420 vannamei, but was shown to play no role in dsRNA-induced antiviral immunity (Labreuche et 421 al., 2009). Wang et al. (2011) reported investigate Tumor necrosis factor receptor (TNFR)-422 423 associated factor 6 (TRAF6) function in invertebrate innate immune responses, Litopenaeus vannamei TRAF6 (LvTRAF6) was identified and characterized. 424

425 Superoxide dismutases (SODs), which are important antioxidant enzymes, are present in almost all oxygen respiring animals. In the present study, cMnSOD was found to have basal 426 427 expression level in shrimp injected with PBS while in shrimp injected with VP28dsRNA, the mRNA expression was found to be increased significantly in haemocytes, gill and 428 429 hepatopancreas. This indicates that dsRNA-VP28 stimulates immune genes reaching maximum level at 12 hp.i., thereafter decreasing gradually after 24 hp.i. and finally reaching normal level at 430 431 48 hp.i. The prophenoloxidase system forms an important part of an immune recognition process 432 of the defense mechanism and enzyme production during proPO system activation. Takahashi et al. (2000) observed increased proPO activity in *M. japonicas* fed with diet containing LPS for 7 433 days at a dose of 20 mg kg⁻¹ shrimp body weight. In *Penaeus monodon* that had been fed a diet 434

containing glucan for 20 days, proPO activity was significantly increased when compared to non 435 glucan group (Chang et al., 2003). In the present study, enhanced mRNA expression was found 436 437 in haemocytes, gill and hepatopancreas in response to injection of dsRNA-VP28 in L. vannamei. This result agrees with previous works carried out by various workers with different 438 immunostimulants. Crustin, one of the important AMPs in crustacean has gained the attention of 439 many researchers. Crustin has been demonstrated to be markedly up regulated in the kuruma 440 prawn, Marsupenaeus japonicus from day 1 to 7 with oral administration of peptidoglycon 441 (Rattanachai et al., 2005). Similarly, increased expression of crustin was found in shrimp 442 injected with dsRNA-VP28at 12 hp.i. and gradually decreased to normal level at 48 hp.i. 443 Immune gene expression profile showed that a yeast incorporated diet up-regulated all the six 444 AMP (antimicrobial peptide genes) genes (ALF, Crustin-1, Crustin-2, Crustin-3, Penaeidin-3 and 445 Penaeidin-5) and immune gene (proPO) to a significant level and that the genes were 446 differentially expressed in the haemocytes of P. monodon during pre-and post-challenge period 447 (Divva et al., 2013). Although the detailed mechanism of proPO, cMnSOD, hemocyanin, crustin, 448 BGBP, LPs, lectin and lysozyme remain unknown, their up-regulation induced by dsRNA-VP28 449 450 indicates the potential effects of dsRNA-VP28 on the defense responses against bacterial and viral diseases. 451

452

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458 **Reference**

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Figure legend

Fig.1 Bacterially expressed dsRNA specific to VP28 gene of white spot syndrome virus. Lane M-100 bp Marker; Lane 1 - Control *E.coli*HT115(DE3) containing LITMUS38i vector withoutinsert; Lane 2 -Un-induced *E. coli*HT115(DE3) containing LITMUS38i with V28 gene; Lane 3 - IPTG induced *E.coli*HT115(DE3) containing LITMUS38i with V28 gene.

Fig.2 Prophenoloxidase activity in haemolymph of *L. vannamei* injected with dsRNA-VP28. Bar represents mean value for ten shrimp with standard error. Asterisk indicates significant difference (*P < 0.05) between control and treated shrimp.

Fig.3 Superoxide dismutase activity in haemolymph of *L. vannamei* injected with dsRNA-VP28. Bar represents mean value for ten shrimp with standard error. Asterisk indicates significant difference (*P < 0.05) between control and treated shrimp.

Fig.4 Superoxide anion level in haemolymph of *L. vannamei* injected with dsRNA-VP28. Bar represents mean value for ten shrimp with standard error. Asterisk indicates significant difference (*P < 0.05) between control and treated shrimp.

Fig.5 Malondialdehyde content in haemolymph of *L. vannamei* injected with dsRNA-VP28. Bar represents mean value for ten shrimp with standard error. Asterisk indicates significant difference (*P < 0.05) between control and treated shrimp.

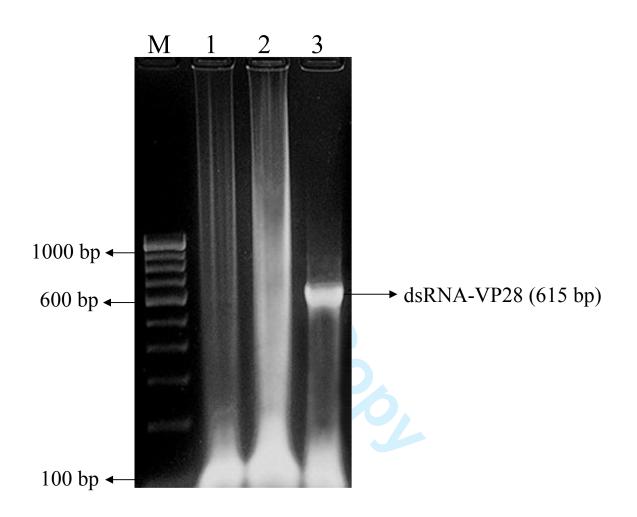
Fig.6 Clotting time (in seconds) of haemolymph from of *L. vannamei* injected with dsRNA-VP28.

Fig.7 Total hemocyte count in L. vannamei injected with dsRNA-VP28 and normal shrimp.

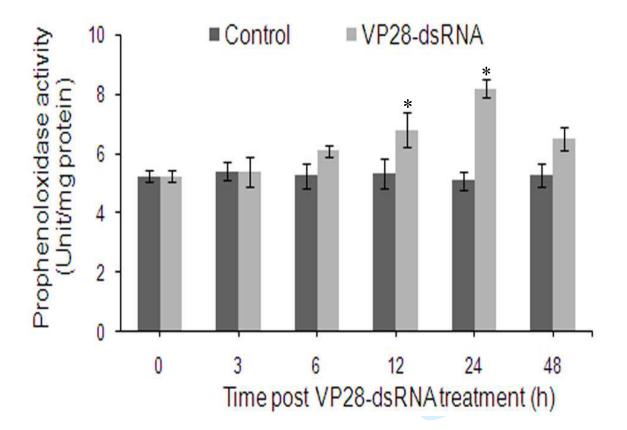
Fig.8 Oxyhemocyanin level in haemolymph of *L. vannamei* injected with dsRNA-VP28 and normal shrimp.

Fig.9 The relative expression level of proPO (A), SOD (B), hemocyanin (C), crustin (D), BGBP (E), lipopolysaccharides (F), lectin (G), lysozyme (H), Toll receptor (I) and Tumor receptor (J) genes in haemocytes, gill and hepatopancreas of *L. vannamei* injected with dsRNA-VP28. Bar represents mean value for ten shrimp with standard error. Asterisk indicates significant difference (*P < 0.05) between control and treated shrimp.

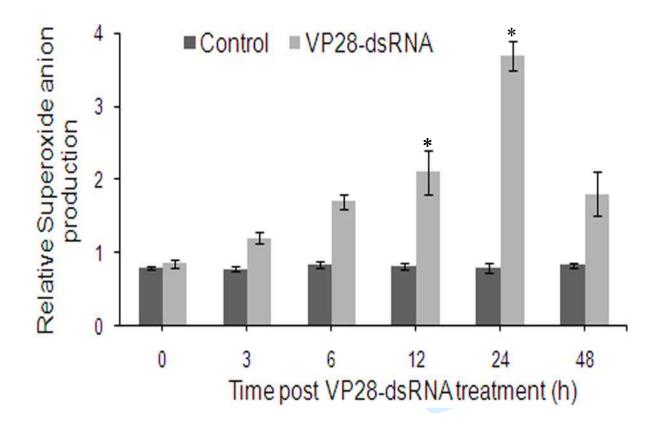
Fig.1













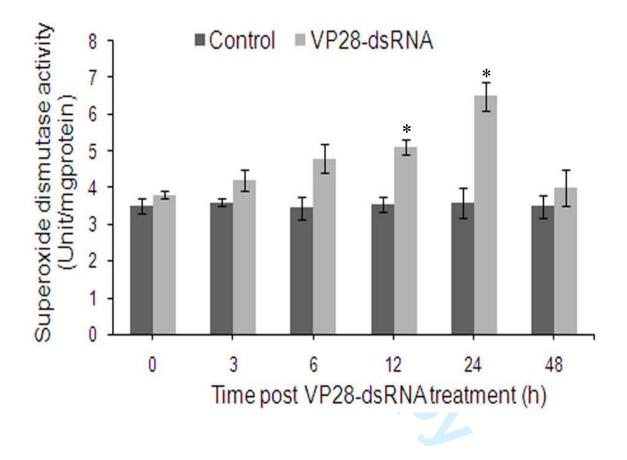


Fig.5

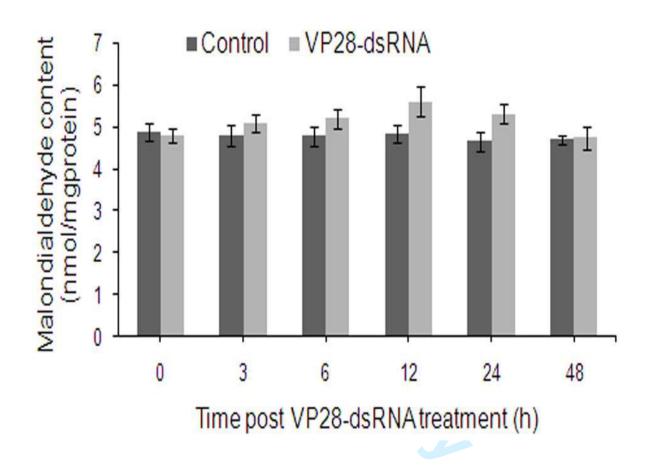


Fig.6

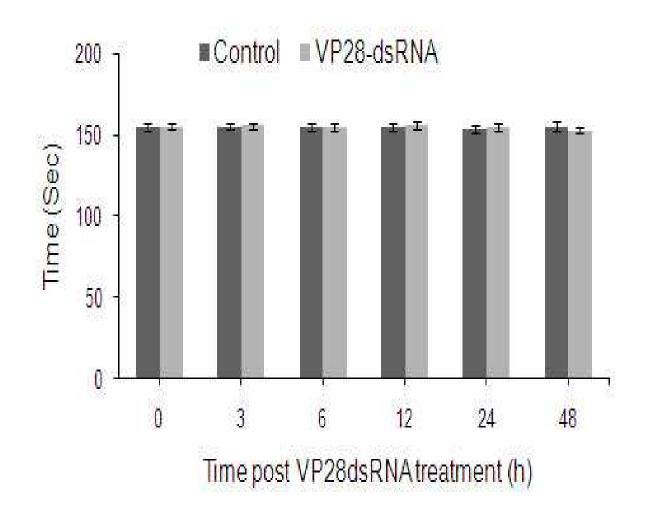
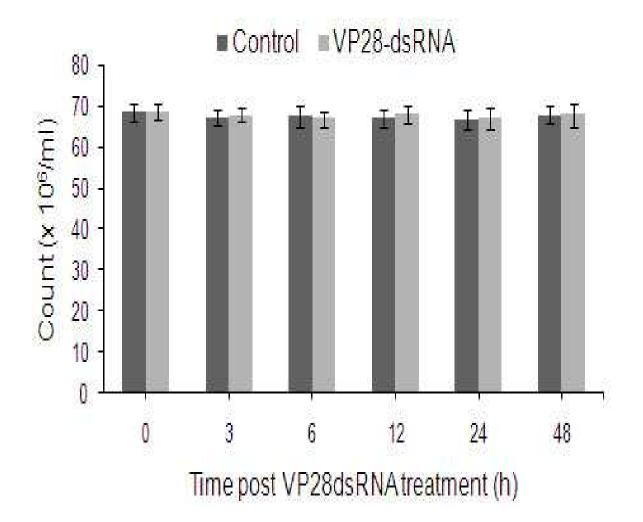


Fig.7





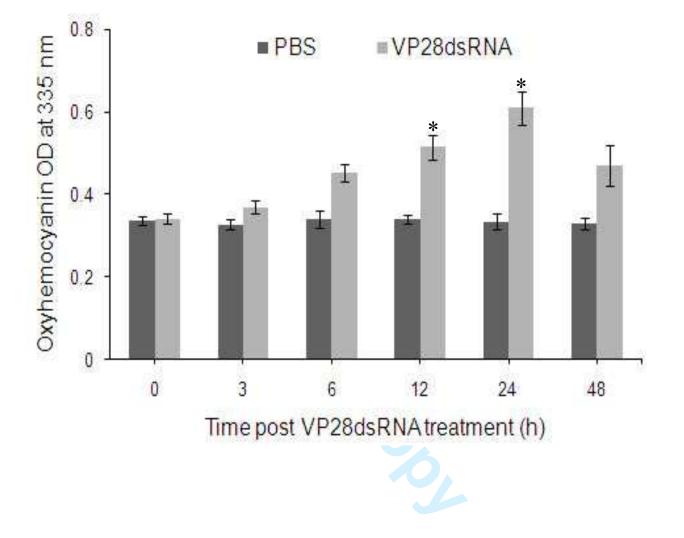
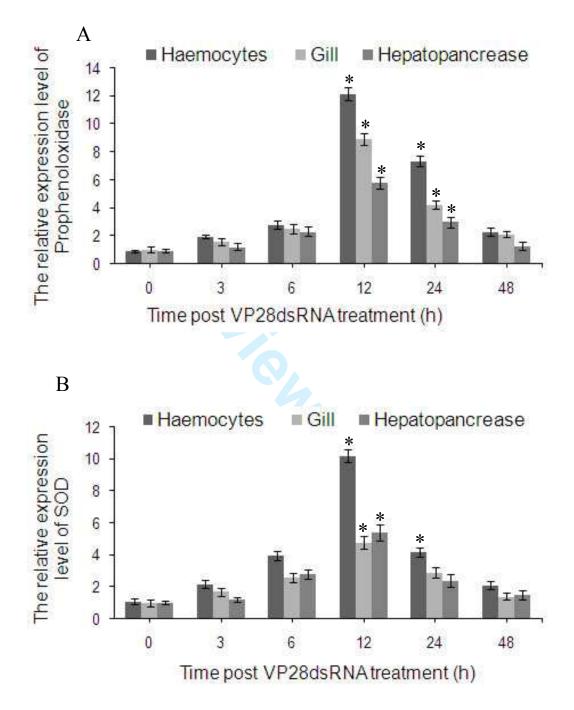
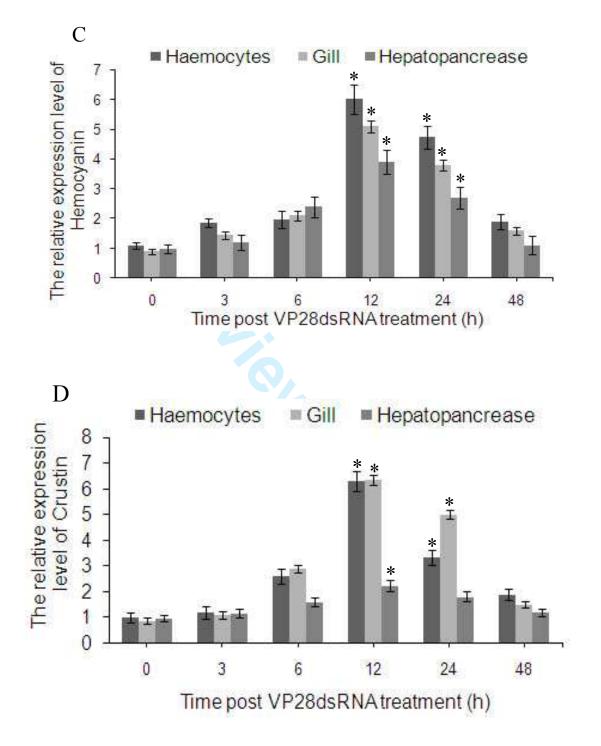
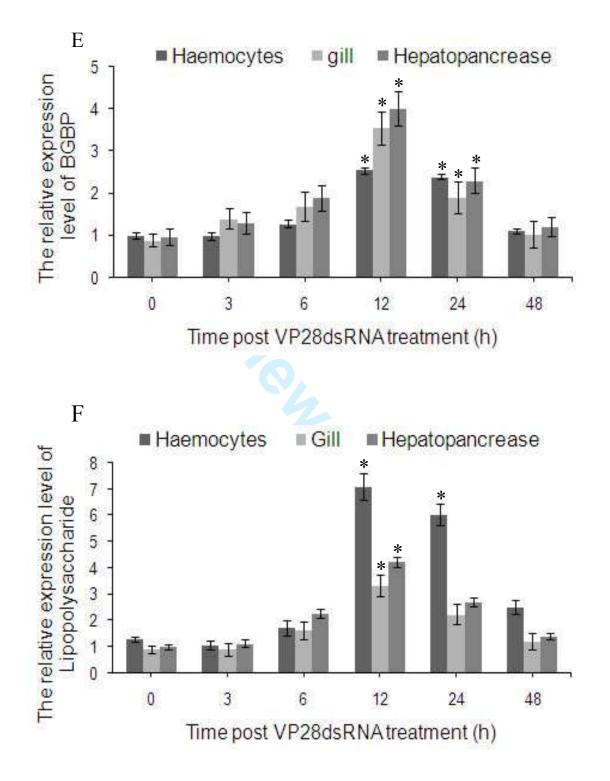
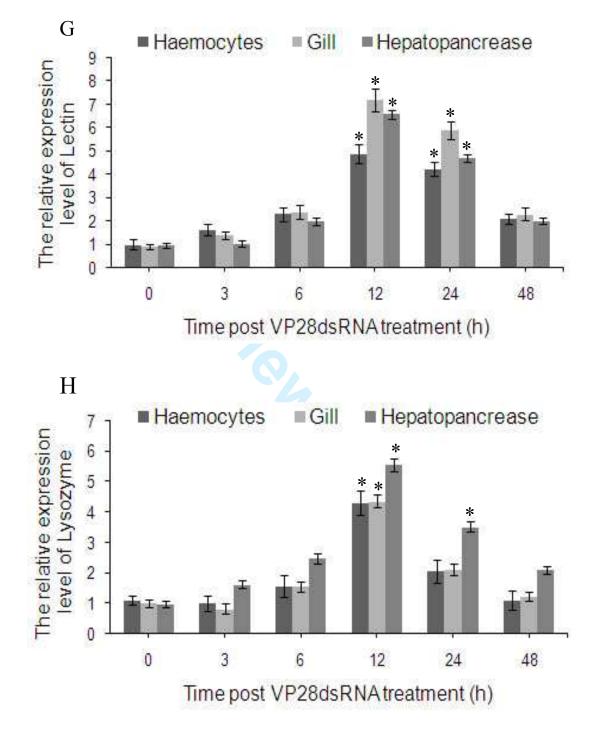


Fig.9









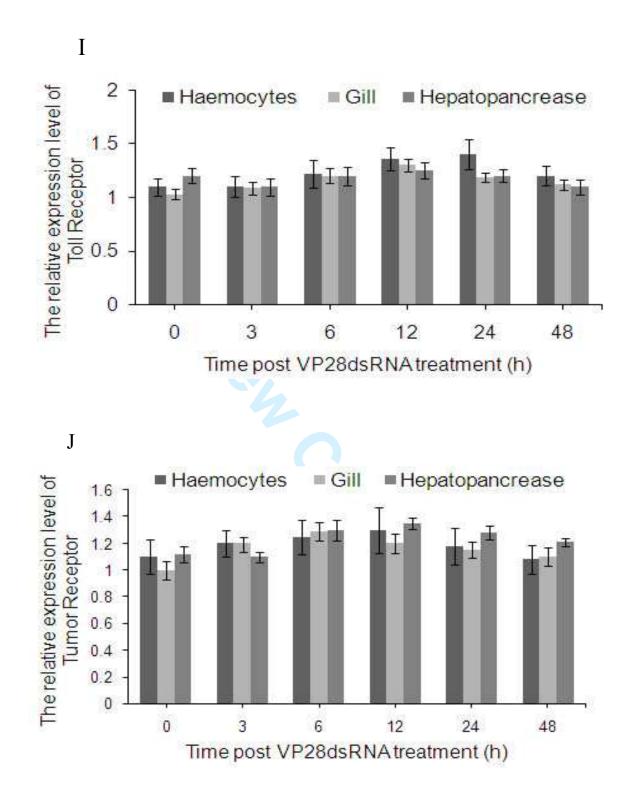


Table 1 Primers and	nd their nucleotide sequences	used immune genes	in Litopenaeus vannam	<i>iei</i> in the
present stu	udy.			

Primers	Sequence (5'-3')	GenBank	
FILLEIS	Sequence (5 - 5)	Accession No.	
β-Actin F	AGTAGCCGCCCTGGTTGTAGAC		
β-Actin R	TTCTCCATGTCGTCCCAGT	AF300705.2	
Prophenoloxidase F	CTCCATTCCGTCCGTCTG		
Prophenoloxidase R	CGGCTTCGCTCTGGTTAG	AY723296.1	
cMnSOD F	CGCGGATCCGATGGCTGAGGCAAAGGAA		
cMnSOD R	CCGGAATTCTGGGCAAACATCTGTGCTATCT	DQ005531.1	
Crustin F	GGCGGAGTAGGTGTTGGT	AY486426.1	
Crustin R	R TGTGGGCAGTCGAGTATCTT		
Lipopolysacharides F	CATGTCCAACTTCGCTTTCAGA	Character 1 2005	
Lipopolysacharides R	ATCACCGCGTGGCATCTT	Chen et al., 2005	
Lysozyme F	CGCGGATCCGACTGATGCGGAAGCGACTA	AY170126.2	
Lysozyme R	CCGGAATTCAGCCACCCAGGCAGAATA	AY1/0120.2	
Lectin F	TGTCGCCAGTGTTCGTTC		
Lectin R	CTACTTCATCTGTTGCTCCT	EF583939.1	
BGBP F	TACGGCTGCTCCCGAACT		
BGBP R	TACGAGGCAACATCGAAATA	AY249858.1	
Toll Receptor F	CTATTGTGGTGCTTTCGT		
Toll Receptor R	TCTCGTCCAACTCGCTCT	DQ923424.1	
Tumor Receptor F	CGCGGATCCGATGTTCACCGACACCTAT		
Tumor Receptor R	CCGGAATTCTTCCAACTTTGCCTTTAC	HM581680.1	
Hemocyanin F	TCCTCGTCTCAACCACAA	HQ709161.1	
Hemocyanin R	anin R GCACATCAGGCACAGTAA		

Table 2 Quantification of mRNA expression of immune-related genes in haemocytes, hepatopancreas and gill of shrimp injected with dsRNA-VP28 by real time PCR. Mean (\pm SE) values of cycle threshold (C_T)

	Cycle threshol			plicate assays for		n from normal
Immune genes		and VP2	28dsRNA inject	ed shrimp (L. val	nnamei)	
	Normal (0 h)	3 h	6 h	12 h	24 h	48 h
Crustin						
Haemocytes	31.77±0.78	30.95±0.27	29.38±0.14	27.38±1.20*	28.21±0.10*	30.44 ± 0.90
Gill	32.45±0.33	32.05±0.61	30.29±0.28	28.90±1.62*	29.15±0.72*	31.87±1.13
Hepatopancreas	30.85±0.30	30.40±0.71	30.10±1.40	29.93±0.52*	30.12±1.03*	33.01±1.22
Lipopolysaccharide						
Haemocytes	22.97±0.06	22.50±0.20	21.73±1.04	18.27±0.22*	18.81±0.97*	21.90±0.36
Gill	26.79±0.19	26.88±0.41	25.99±0.72	24.65±0.92*	25.16±0.54*	26.01±1.35
Hepatopancreas	28.96±0.48	28.50±0.34	27.44±0.19	26.01±0.33*	$27.20{\pm}1.02$	27.98 ± 0.28
Lysozyme						
Haemocytes	21.80±0.02	21.71±0.52	21.09±0.73	20.53±1.41*	20.70 ± 0.20	21.01±1.26
Gill	27.17±0.35	27.54±0.56	26.57±0.33	25.87±0.90*	26.01±1.01*	26.99 ± 0.81
Hepatopancreas	28.49±0.19	27.05±0.12	26.38±0.17	25.01±0.10*	25.42±0.29*	27.30 ± 1.40
Tumor Receptor						
Haemocytes	32.98±0.39	31.92±0.51	30.84 ± 0.54	31.33±0.25	31.10±0.39	31.90 ± 0.82
Gill	31.92±0.41	31.41±0.19	31.02 ± 0.41	30.47±0.34	31.12±0.64	31.57 ± 0.70
Hepatopancreas	31.73±0.22	31.69±0.63	31.32±0.29	31.50±0.92	31.02 ± 0.56	31.20 ± 0.80
Lectin						
Haemocytes	32.55±1.24	31.10±0.12	30.89±0.72	29.34±0.22*	29.70±0.23*	31.90 ± 1.04
Gill	31.76±0.45	30.78 ± 0.74	30.01±0.29	27.56±0.17*	27.69±0.10*	30.97 ± 0.58
Hepatopancreas	29.15±0.46	28.01±1.01	27.87 ± 0.48	25.19±0.32*	25.31±0.10*	28.89 ± 0.61
BGBP						
Haemocytes	29.23±0.18	30.66±0.31	29.01±0.11	27.62±1.57*	27.90±0.20*	28.75±0.29
Gill	31.10±0.64	31.06±0.76	30.76±0.65	28.77±0.70*	29.36±0.71*	30.95±0.93
Hepatopancreas	28.70±0.18	28.31±0.74	28.04±0.18	27.44±0.29*	27.79±0.48*	28.20 ± 0.80

Table 2 Continu	ed
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Toll Receptor						
Haemocytes	25.75±0.46	25.13±0.30	24.42 ± 0.38	24.65±0.81	24.13±0.17	24.90±1.20
Gill	30.88±0.13	30.59±0.46	30.10±0.27	29.55±0.63	29.13±0.99	29.96±0.30
Hepatopancreas	32.80±0.14	32.10±0.61	31.95 ± 0.10	31.30±0.60	31.18±0.12	31.83±0.38
Prophenoloxidase						
Haemocytes	30.84±0.27	29.89±0.25	29.22 ± 0.80	25.18±0.22*	26.65±0.84*	29.79±1.03
Gill	32.51±0.17	31.47±0.50	30.12±0.91	27.10±0.47*	29.96±0.22*	31.93±1.25
Hepatopancreas	31.53±0.37	30.75±0.61	30.95 ± 1.08	28.44±0.40*	29.10±0.79	30.66±0.19
Hemocyanin						
Haemocytes	30.99±0.51	30.11±0.30	29.35±0.14	27.76±0.65*	28.01±0.26*	30.20 ± 0.80
Gill	29.77±0.24	29.40±0.14	28.90±0.61	26.34±0.44*	28.80±0.75*	29.10±0.24
Hepatopancreas	28.98 ± 0.05	28.66±0.19	28.14±0.37	27.10±0.14*	27.50±0.16*	28.42 ± 0.28
SOD						
Haemocytes	33.84±0.20	32.68±0.23	31.98±0.63	29.12±0.10*	31.68±0.53	33.41±0.82
Gill	31.98±0.43	31.55±0.09	31.12±0.12	30.02±0.97*	31.50±0.18	31.88±0.64
Hepatopancreas	32.05±0.36	31.80±0.15	30.96±0.83	30.15±0.42*	31.11±0.35	31.90 ± 0.40
β-Actin						
Haemocytes	27.33±1.20	27.56±0.79	27.95 ± 1.20	27.82±0.19	28.15±0.71	28.01±0.95
Gill	28.01±0.43	28.12±0.48	28.10 ± 0.70	28.08±0.71	28.90±0.50	29.01±0.20
Hepatopancreas	28.89 ± 0.52	29.01±1.02	29.15±0.91	28.98±0.18	28.88±0.27	28.40 ± 0.70