


Article

Dietary β -1,3-Glucan Promotes Growth Performance and Enhances Non-Specific Immunity by Modulating Pattern Recognition Receptors in Juvenile Oriental River Prawn (*Macrobrachium nipponense*)

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Abstract: As a typical pathogen-associated molecular pattern (PAMP), β -1,3-glucan can engage with pattern recognition receptors (PRRs) to initiate an immune response. In this study, we investigated the effects of dietary β -1,3-glucan on growth performance, antioxidant capacity, immune response, intestinal health, and bacterial resistance in juvenile *Macrobrachium nipponense*. Prawns were fed with five experimental diets containing 0%, 0.05%, 0.1%, 0.2%, and 0.4% β -1,3-glucan for eight weeks. The findings demonstrated that the inclusion of β -1,3-glucan improved weight gain and survival rate in prawns. Prawns fed with β -1,3-glucan exhibited elevated activities of hepatopancreatic ACP (acid phosphatase), AKP (alkaline phosphatase), and SOD (superoxide dismutase), while MDA (malondialdehyde) content was reduced. Expression levels of PRRs related genes including *LGBP* (lipopolysaccharide and β -1,3-glucan binding protein), *lectin*, and *LBP* (lipopolysaccharide-binding protein) were significantly increased in prawns fed with β -1,3-glucan. Intestinal flora analysis revealed suppression of Cyanobacteria abundance at the Phylum level and enhancement in *Rhodobacter* abundance at the genus level in prawns fed with a 0.2% β -1,3-glucan diet. Furthermore, prawns fed with 0.1%, 0.2%, and 0.4% β -1,3-glucan demonstrated significantly higher survival rates following *Aeromonas hydrophila* infection. In conclusion, β -1,3-glucan can activate PRRs to improve immune responses in *M. nipponense*. Within the range of β -1,3-glucan concentrations set in this experiment, it is recommended to add 0.18% of β -1,3-glucan to the diet, taking into account the positive effect of β -1,3-glucan on the survival rate of *M. nipponense*.

Keywords: *Macrobrachium nipponense*; β -1,3-glucan; PRRs; non-specific immunity; intestinal flora; *Aeromonas hydrophila*

Key Contribution: The potential mechanism of β -1,3-glucan in enhancing the immune capacity of *M. nipponense* was investigated with a focus on pattern recognition protein receptors and intestinal health. The optimal levels of β -1,3-glucan supplementation were determined based on survival rates.



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1. Introduction

The aquaculture industry is experiencing rapid global growth and it is recognized as the fastest-growing sector in worldwide food production, accompanied by a surge in demand for fish products in the consumer market [1]. Ensuring the sustainable health of aquatic organisms is an important endeavor for the human food supply chain. Disease outbreaks are widely acknowledged as a major impediment to the growth of the aquaculture industry, resulting in substantial global losses annually [2]. Antibiotics are now widely used in the aquaculture industry to minimize losses of aquatic products caused by

bacterial pathogens [3]. Yet, antibiotic-resistant (AMR) bacteria are also born out of the feed industry's practice of indiscriminate and constant abuse of production [4], its genes and residues come with it [5]. When AMR bacteria evolve and spread in natural water bodies, they not only cause disease and death in aquatic animals but also have a profoundly negative impact on economic development [6]. Leading some countries to impose restrictions or bans on the widespread antimicrobial agent usage in aquaculture [7]. Consequently, there is an imperative need for non-toxic, environmentally friendly, and highly efficient biological agents that can serve as alternatives to antibiotics within the aquaculture industry. Immunostimulants represent one such option that can effectively mitigate the outbreak of aquatic animal diseases [8]. Understanding natural immune responses in crustaceans plays an important role in studies applying immunotherapy to enhance disease resistance among cultured animals.

Crustaceans are known to lack an acquired immune system, relying mainly on innate immunity [9]. PRRs (pattern recognition receptors) enable the immune system to distinguish between "self" and not-self", playing a crucial role in initiating the immune response [10]. Various PRRs have been established in crustaceans, including *LBPs* (lipopolysaccharide-binding proteins) [11], *LGBPs* (lipopolysaccharide and β -1,3-glucan binding proteins) [12], *GNBPs* (gram-negative binding proteins) [13], *PGRPs* (peptidoglycan recognition proteins binding proteins) [14], *bGBPs* (β -glucan binding proteins) [15], *CTLs* (C-type lectins) [16], *TEPs* (sulfate-containing proteins) [17], *SCRs* (scavenger receptors) [18], *FBGLs* (fibrinogen-like structural domain immunoselectors) [19], and *TLRs* (Toll-like receptors) [20]. Upon recognition of corresponding PAMPs, PRR-PAMP interactions trigger a cascade of immune reactions. Phagocytosis mediated primarily by hyalinocytes eliminates foreign invasive substances [21]; primarily encapsulation and cytotoxicity predominantly carried out by semigranular cells counteract parasites and fungi invading the host [22]; activation of Toll and IMD (immune deficiency) signaling pathways regulates antimicrobial peptides production [23,24]; *lectin* secretion and proPO-AS (Prophenoloxidase activating system) activation occur as well [25,26]. Furthermore, we observed a correlation between immunization and growth performance in crustaceans [27–29]. Therefore, it is worthwhile to explore the research directions that stimulate the interaction between PRRs and PAMPs using immunostimulants to enhance environmental resistance and growth performance in crustaceans.

Immunostimulants are effective in boosting non-specific immune responses and improving overall immune system function [30]. Numerous studies have demonstrated that incorporating immunostimulants into feeds leads to improved immunity in crustaceans [31–33]. One prominent immunostimulant in crustaceans is β -1,3-glucans, a carbohydrate group synthesized by the cell walls of plants, bacteria, fungi, and some freshwater algae. These glucans are well-sourced and inexpensive [34]. They can be either branched or linear polymers (polysaccharides) composed of repeating glucose units linked by β -glycosidic bonds. Remarkably, previous studies have shown that β -1,3-glucan acts as a typical PAMP interacting with PRRs present in immune cells to trigger an innate immune response [35,36]. As a dietary additive, β -1,3-glucan is widely used in aquatic animals to bolster immunity, enhance disease resistance, and stimulate growth [37]. For instance, feeding juvenile tiger shrimp (*Penaeus esculentus*) with diets containing β -glucan and mannan oligosaccharide improves immune response and survival when exposed to WSSV (*white Spot Syndrome Virus*) infection [38], while feeding *M. nipponense* diets enriched with β -1,3-glucan not only improves its innate immunity and growth performance but also modulates the diversity of gut microbiota for maintaining gut health [39,40]. In another study involving *Eriocheir sinensis* fed β -1,3-glucan followed by *Vibrio parahaemolyticus* infection after knocking out *LGBP* at the entry point of PRRs for β -1,3-glucan resulted in increased mortality rates [41]. Similarly, higher expression levels of *LGBP* were detected in the hepatopancreas of *Litopenaeus vannamei* fed a diet supplemented with β -glucan [42]. These findings highlight the prominent research interest surrounding the use of β -1,3-glucans as trace additives in aquaculture. Therefore, β -1,3-glucan represents a promising

micro-additive for immune-enhancing feeds with the potential to reduce and replace the dependence on antibiotics in the aquaculture industry.

In China and other Asian countries, the oriental river prawn, *M. nipponense*, is widely distributed in freshwater basins. Specifically in China, the annual production of this prawn is 240,739 tons, making it one of the most commercially viable species [43–45]. However, intensive prawn farming in *M. nipponense* has led to deteriorating water quality and environmental stress, making prawns susceptible to bacterial diseases, mass death events caused by *Aeromonas veronii* [46], and “red gill disease” from *M. nipponense* [47]. Due to multiple adverse effects associated with antibiotic usage [48], many countries have prohibited their use in aquatic organisms intended for human consumption like fish and shrimp due to concerns over negative impacts on human health, development of antibiotic-resistant strains, reduced efficacy from repeated applications, and environmental contamination issues [49–51]. Consequently, finding viable alternatives that are effective yet somewhat economical and environmentally safe has become an urgent requirement for restraining antibiotic usage in *M. nipponense* farming practices. This study sought to evaluate the impacts of β -1,3-glucan diets on growth capability, antioxidant power, immune response, digestive well-being, and immunity against *M. nipponense* bacterial infections. Practical insight into alternative treatment options for ensuring sustainability within *M. nipponense* aquaculture is provided by the results obtained.

2. Materials and Methods

2.1. The Origin of Prawns and β -1,3-Glucan

The prawns selected in this experiment were healthy, with similar growth age, and no obvious scar. The prawns were all supplied by Wuyue Agricultural Co., Ltd. (Huzhou, China). The β -1,3-glucan utilized in the study was obtained from Nanjing Taixin Biotechnology Co., Ltd. (Nanjing, China) with a purity of 95.21%.

2.2. Experimental Design and Dietary Composition

The basal diet was made as an isonitrogenous and isocaloric diet using fishmeal, soybean meal, and rapeseed meal as protein sources, and fish oil and soybean oil as lipid sources. The experimental diets were prepared by supplementing the basic diet with 0% (control), 0.05%, 0.1%, 0.2%, and 0.4% β -1,3-glucan, referred to as B0, B1, B2, B3, and B4, respectively (Table 1).

In order to create experimental diets, all ingredients need to be processed individually. The initial step involved grinding the larger particles of the ingredients. Subsequently, a more homogeneous mixture of particles and powders was obtained using a 212 μ m sieve screen. Following this, it was ensured that the individually bagged powders exhibited uniform and consistent quality. The individual ingredients were accurately weighed according to the list of diet ingredients (Table 1) using a 0.1 mg electronic scale. The finished weighed solid ingredients were added to the mixing vessel with accurately weighed semi-solid and liquid ingredients (fish oil, soybean oil, and distilled water). The mix was produced in preliminary strip form using a twin-screw extruder (School of Chemical Engineering, South China University of Technology, Guangzhou, China) and finally dried in a forced-air oven at 40 °C. The dried feeds were stored in sealed bags at –20 °C.

The determination of moisture, crude lipid, and crude protein contents in each experimental diet was conducted by using standard procedures [52]. To determine moisture, the sample was dried at 105 °C for 24 h to a constant weight. The Dumas nitrogen determination apparatus combustion method was utilized to determine the crude protein content. The ash content was determined by setting the parameter 550 °C in a Muffle furnace for 6 h and Soxhlet extraction method was employed to measure total lipid (Table 1).

Table 1. Ingredients and proximate composition of basal diets (% dry weight).

Ingredients	B0	B1	B2	B3	B4
Fish meal	18	18	18	18	18
Soybean meal	40	40	40	40	40
Rapeseed meal	15	15	15	15	15
Fish oil	3	3	3	3	3
Soybean oil	1.5	1.5	1.5	1.5	1.5
Corn starch	10	10	10	10	10
Soy lecithin	0.5	0.5	0.5	0.5	0.5
Cholesterol	0.5	0.5	0.5	0.5	0.5
Vitamin premix ^a	2	2	2	2	2
Mineral premix ^b	3	3	3	3	3
Choline chloride	0.5	0.5	0.5	0.5	0.5
Sodium carboxymethyl cellulose	2	2	2	2	2
Cellulose	4.00	3.95	3.90	3.80	3.60
β -1,3-glucan ^c	0	0.05	0.10	0.20	0.40
Total	100	100	100	100	100
Proximate composition					
Dry matter	91.52	91.91	91.72	92.19	91.80
Crude protein	38.44	38.56	38.44	38.46	38.36
Crude lipid	9.73	9.77	9.78	9.70	9.78
Ash content	9.32	9.60	9.42	9.55	9.31

^a Vitamin mixture (per kg mixture): vitamin A 4,200,000 IU, vitamin C 60,000 mg, α -tocopherol acetate 20,000 mg, vitamin D3 1,200,000 IU, vitamin K 10,000 mg, vitamin B1 7500 mg, vitamin B2 1000 mg, vitamin B6 16 mg, vitamin B12 20 mg, nicotinic acid 50,000 mg, folic acid 4000 mg, inositol 60,000 mg, biotin 100 mg, calcium pantothenate 35,000 mg. ^b Composition of mineral mixture (per kg): KCl 28 g, MgSO₄·7H₂O 100 g, NaH₂PO₄ 215 g, KH₂PO₄ 100 g, Ca (H₂PO₄)₂·H₂O 265 g, CaCO₃ 105 g, C₆H₁₀CaO₆·5H₂O 165 g, FeC₆H₅O₇·5H₂O 12 g, ZnSO₄·7H₂O 4.76 g, MnSO₄·H₂O 1.07 g, AlCl₃·6H₂O 0.15 g, CuCl₂·2H₂O 0.24 g, CoCl₂·6H₂O 1.4 g, KI 0.23 g, α -cellulose 2.15 g. ^c Provided by Nanjing Taixin Biotechnology Co., Ltd., Nanjing, China.

2.3. Experimental Animals and Feeding Trial

At the outset, the tanks used for the experiment were sterilized and dried. The prawns were acclimatized for seven days in a pre-raised environment. During this period, a commercial feed with 39% protein was consumed by them. Following the adaptation period, fifteen tanks (each with a volume of 300 L) were randomly chosen to stock uniformly sized, healthy prawns with an initial average weight of 0.10 g, with each tank housing seventy prawns. The prawns were fed twice a day at 8:00 a.m. and 5:00 p.m., depending on their average weight (about 4–6% of the body weight). They were fed β -1,3-glucan diets for eight weeks. Throughout the eight-week experimental period, it was ensured that the daily water exchange was one-third of the original volume. After the water exchange was completed, water quality parameters were monitored in real-time to maintain a temperature between 25 and 28 °C, a total ammonia and nitrate content of less than 0.1 mg/L, and a dissolved oxygen content in the water of more than 6.5 mg/L, with the following parameters being maintained. The cycle of light and darkness under natural daylight conditions is fourteen hours of light and ten hours of darkness.

2.4. Analysis of Growth-Related Parameters

At the end of the eight-week feeding experiment, the entire tank stock of prawns was counted and weighed after 24 h of cessation of feeding, and the final body weights (FBW) and initial body weights (IBW) obtained were used to calculate the final growth performance involving the following equations:

$$\text{WG (\%)} = 100 \times (\text{Final weight} - \text{Initial weight}) / \text{Initial weight}$$

$$\text{SGR (\%/d)} = 100 \times (\text{Ln Final weight} - \text{Ln Initial weight}) / \text{breeding days}$$

$$\text{Survival rate (\%)} = 100 \times \text{mantissa of final prawns} / \text{mantissa of prawns}$$

2.5. *Aeromonas Hydrophila* Challenge

The strain of *A. hydrophila* used in this experiment was provided by Bei Na Biological Company. For the *A. hydrophila* challenge test, 12 prawns were randomly selected from each tank in the experimental group. The use of *A. hydrophila* solution at a concentration of 1.2×10^7 CFU/mL in a dose of 20 μ L was determined based on the IFW and takedown pre-tests as a takedown reagent, and this reagent was injected into the cartilage of the basal joint of the prawns of the third maxillipeds. The cumulative survival rate of the prawns in each tank was recorded every three hours for a continuous period of three days (72 h) following injection with *A. hydrophila*.

2.6. Biochemical Parameters Analysis

Biochemical parameter analysis was performed by randomly sampling 10 prawns from each tank. Each prawn was anesthetized by immersion in 4 °C water. Subsequently, the hepatopancreas was dissected at the cephalothorax for sampling purposes. The ventral sinus was used to collect individual hemolymph and stored at –80 °C for future analysis. Prawn hepatopancreas samples were soaked in 0.86% saline and centrifuged at $1500 \times g$ for twenty minutes at 4 °C. The enzymatic activities of specific biochemical enzymes were analyzed using hepatopancreas samples. The superoxide dismutase (SOD) activity assay kit was purchased from Suzhou Comin Biotechnology Co., Ltd. (Suzhou, China, SOD-2-W). The kits (Nanjing Jianjian Bioengineering Institute, Nanjing, China) were used to detect acid phosphatase (ACP, A060-1-1), alkaline phosphatase (AKP, A059-1-1) activities, and malondialdehyde (MDA, A003-1-1) activities in hepatopancreas and serum. A full-wavelength microplate reader (Thermo Scientific Multiskan GO 1510, Shanghai, China) was used to measure absorptions during analysis procedures. The other part of the hepatopancreas sample was kept for the sole purpose of extracting total RNA (ribonucleic acid) for analyzing mRNA (messenger ribonucleic acid) expression.

2.7. 16S Illumina High-Throughput Sequencing Analysis Process

Intestinal flora and growth performance are often closely linked. In order to explore the effect of different levels of β -1,3-glucan (low, medium, and high) on prawns, 15 prawns from each tank in B0, B2, and B4 were selected as a representative reference for measuring intestinal flora. A TIANamp Micro DNA Purification Kit (Tiangen, Beijing, China, GDP316) was used to isolate the intestinal contents of ten prawns and extract the total bacterial community DNA. DNA samples were analyzed using a NanoDrop spectrophotometer (Thermo Fisher Scientific, Waltham, MA, USA) for their quality and quantity. In the following step, the DNA samples were mailed to Origingene Biological Medicine Technology Co. (Shanghai, China) for 16S intestinal flora analysis.

Sequence data from the V3-V4 region of the bacterial 16S rRNA gene were amplified with prokaryotic primers F341/R806. Each primer had unique 8-bp barcodes added to distinguish the products of polymerase chain reaction (PCR). A different 8bp barcode is added to each primer in order to differentiate the final product of the PCR. FastPfu Buffer, FastPfu Polymerase, primer, dNTPs, and template DNA are added and mixed proportionally in the reaction system, and amplification was accomplished under the following PCR setup program: denaturation of DNA at 95 °C for 5 min, followed by denaturation at 95 °C for 30 s, annealing at 55 °C for 30 s, extension at 72 °C for 45 s, and finally extension at 72 °C for 10 min, for a total of 27 cycles. Illumina high-throughput sequencing allows purification of PCR products, the results of which have been accepted at GenBank, and since the former experiment was submitted together with the samples from this experiment, the package can now be accessed via <https://www.ncbi.nlm.nih.gov/bioproject/954951> (accessed on 12 April 2023) and the SRA search number PRJNA954951 expanded. Quality-filtered raw reads using QIIME (version 1.17) analysis were carried out. The reads were clustered in high quality using USEARCH version 7.1, with a similarity threshold of 97% as the basis for generating OUT (Operational Units of Classification). The SILVA rRNA database was used as data support and the 16S rRNA sequences were

correlated and analyzed using the RDP Classifier, which sets a confidence threshold of 70%. Finally, community diversity and species richness estimates were created in Mothur in version v.1.30.1.

2.8. mRNA Expression Analysis

The trizol method was used to extract the total RNA from the hepatopancreas samples of 5 prawns from each tank. The Thermo NanoDrop 2000 nucleic acid and protein analyzer (Thermo Fisher Scientific, Waltham, MA, USA) was used to determine the concentration and purity of RNA. Total RNA was reverse transcribed to cDNA under the instructions of the Reverse Transcription Kit (Takara, Japan) and stored in a refrigerator at $-20\text{ }^{\circ}\text{C}$ for future use. Quantitative real-time polymerase chain reaction (qRT-PCR) was used to examine the mRNA expression of immune-related genes in the hepatopancreas. The analyzed genes include lipopolysaccharide and *LGBP*, *lectin*, *LBP*, tumor *p53*, and cysteine-aspartic acid protease 3 (*caspase 3*). The mixture reaction system was 20 L, including 7.6 μL H_2O , primers (10 μM) of 0.2 L each, 10 μL $2\times$ SYBR Green Premix Ex Taq, and 2 μL cDNA. The qRT-PCR instruments were as follows: 10 min at $95\text{ }^{\circ}\text{C}$, followed by 40 cycles of 10 s at $94\text{ }^{\circ}\text{C}$, 30 s at $58\text{ }^{\circ}\text{C}$, and 32 s at $72\text{ }^{\circ}\text{C}$. The mixture reaction system was used to generate the appropriate melting curves after the qRT-PCR reaction to determine the correctness of the results. Three replicate assays were completed each time including a negative control lacking cDNA, and amplification efficiencies were guaranteed to be between 95% and 103%, with correlation coefficients greater than 0.98 for each gene. 18S ribosomal RNA (18S rRNA) was used as an internal reference gene. The primer sequences of the genes used can be seen in Table 2. Gene mRNA expression was calculated using the $2^{-\Delta\Delta\text{CT}}$ [53] quantification method. The cDNA sequences of the oriental river prawn from Genbank were used to design the primers and then synthesized by Sangon Biotech Co., Ltd. (Shanghai, China).

Table 2. Primers used in this study.

Primer	Sequence (5'→3')	GenBank	Product
(Mn) 18S rRNA-S (Mn) 18S rRNA-A	TACTGCTGAGCCGAAGAT CCACGGACTATTACTACCTAC	EU118285.1	159
(Mn) lectin-S (Mn) lectin-A	AAGGGCAAGGTGTCTCTTCG CCTCCCATGGTGTCCATGTC	PP516428	160
(Mn) LBP-S (Mn) LBP-A	GTCTGTCTAGCAAGGGCGTT AGTGTGATGCGATGAGCGA	PP516429	159
(Mn) LGBP-S (Mn) LGBP-A	CTGCTGATATCGTCGACCCC GGCATAGCTGATGCTACGGT	AGF86400.1	167
(Mn) P53-S (Mn) P53-A	TGCTTGCTCACAGCGATAAACTT AGTCGCCGAGTGTCAAGTCAATAT	KT963043.1	112
(Mn) caspase 3-S (Mn) caspase 3-A	TTGTCATGCAGTACTTGACTGAAGC CCTCATGGGTTGTGCATCATTATA	KX651496.1	171

2.9. Data Analysis and Processing

The Kolmogorov–Smirnov test was used to determine data normality, while Levene's test was used to test for variance homogeneity. After confirming the parametric assumptions, the effect of different diets that contain β -1,3-glucans on the measured parameters of *M. nipponense* was evaluated using an analysis of variance (ANOVA).

After the prerequisite ANOVA showed significant differences, Tukey's multiple range test was used to compare means between groups. Third orthogonal polynomials were used to analyze the survival rate of bioinfestation experiments to correctly estimate the optimal amount of β -1,3-glucans to be added. The cubic equation in this experiment had the highest R^2 value and the lowest p -value when analyzed as a mapping exercise, so polynomial regression analyses were used to determine the optimal ratio of β -1,3-glucan required for optimal survival. The Mantle–Cox test was employed to evaluate survival rates for prawns

infected with *A. hydrophila*. Statistical analyses were performed using SPSS 25.0 (Chicago, IL, USA). The mean \pm standard deviation (Mean \pm SD) is the way results are expressed.

Alpha diversity indexes including Chao1 reflected community richness while Shannon and Simpson indices represented community diversity levels. Coverage analysis determined whether the identified 16S rRNA gene sequences accounted for a majority presence of bacteria in samples.

3. Results

3.1. Growth Performance of Prawns

Different levels of dietary β -1,3-glucan significantly influenced the growth performance of prawns (Table 3; $p < 0.05$). The results indicate that there were no significant differences in FBW, WG, and SGR among prawns fed B0, B3, and B4 diets ($p > 0.05$), while both B1 and B2 showed higher values compared to B4, with B2 being notably superior to the other groups ($p < 0.05$). Prawns fed on the B2 and B3 diets exhibited an increase in survival rate compared to those fed on the B0, B1, and B4 diets ($p < 0.05$). Overall, prawn FBW, WG, SGR, and SR showed an increasing trend followed by a decline with the increasing β -1,3-glucan supplementation in their diets. Utilizing the SR data for linear regression analysis revealed the optimal amount of dietary β -1,3-glucan for juvenile *M. nipponense* was 0.18% (Figure 1).

Table 3. Growth performance of juvenile *M. nipponense* fed with experimental diets containing different concentrations of β -1,3-glucan for 8 weeks.

Diet	IBW (g)	FBW (g)	WG (%)	SGR (%/day)	SR (%)
B0	0.10 \pm 0.01	0.51 \pm 0.05 ^{ab}	391.00 \pm 47.33 ^{ab}	2.91 \pm 0.17 ^{abc}	58.10 \pm 0.83 ^b
B1	0.10 \pm 0.01	0.57 \pm 0.01 ^{bc}	446.00 \pm 5.20 ^{bc}	3.10 \pm 0.02 ^{bc}	60.48 \pm 2.18 ^b
B2	0.10 \pm 0.01	0.60 \pm 0.01 ^c	480.67 \pm 14.47 ^c	3.20 \pm 0.04 ^c	69.52 \pm 2.97 ^c
B3	0.10 \pm 0.01	0.50 \pm 0.07 ^{ab}	375.52 \pm 78.21 ^{ab}	2.84 \pm 0.28 ^{ab}	69.05 \pm 0.82 ^c
B4	0.10 \pm 0.01	0.46 \pm 0.04 ^a	341.62 \pm 42.74 ^a	2.72 \pm 0.17 ^a	48.09 \pm 6.75 ^a

IBW: initial body weight, FBW: final body weight, WG: weight gain, SGR: specific growth rate, SR: survival rate. The value is the average of the three replicates for each treatment \pm SD. The presence of different superscript letters in the same column indicates significant differences ($p < 0.05$).

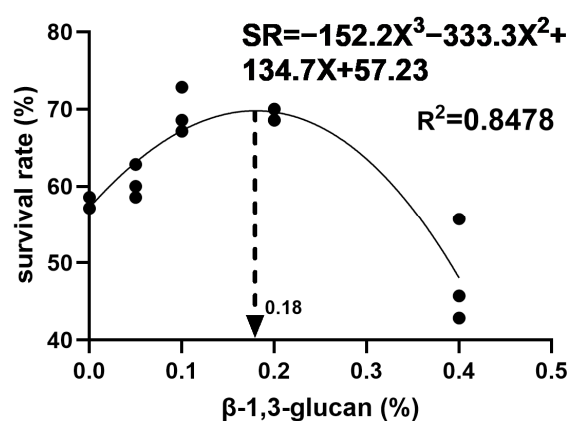


Figure 1. Polynomial regression analysis on survival rate to estimate the optimal β -1,3-glucan supplementation levels in the diet of oriental river prawns (*M. nipponense*).

3.2. Survival Rate of the Prawns after *A. hydrophila* Challenge Test

The cumulative survival rates of *M. nipponense* over a 72 h period are depicted in Figure 2 prawn mortality was primarily observed within the first 12 h, with only isolated individual deaths occurring after 24 h, specifically in groups B2 and B3. The descending order of survival rates for each group was as follows: B2 (79%), B3 (76%), B4 (69%), B1 (63%), and B0 (49%). Notably, groups B2, B3, and B4 exhibited higher survival rates compared to group B0 ($p < 0.05$).

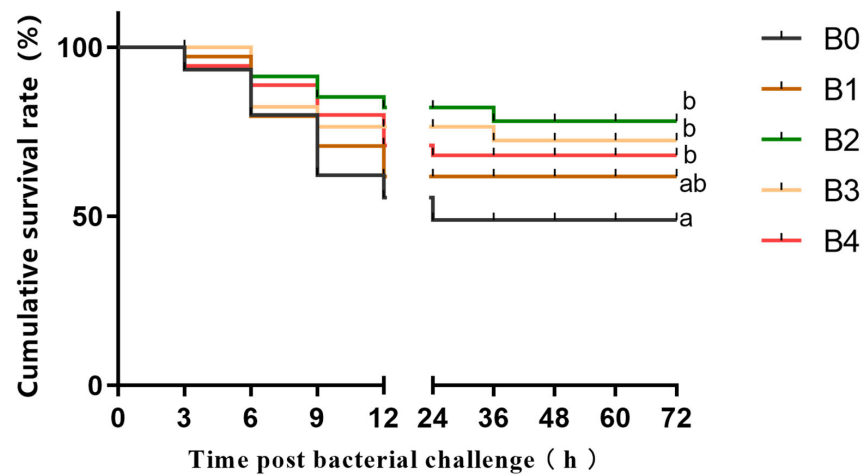


Figure 2. Survival rate of juvenile *M. nipponense* infected with *A. hydrophila* for 72 h. Different colors represent varying concentrations of β -1,3-glucan. The presence of different letters in the figure indicates significant differences ($p < 0.05$).

3.3. Innate Immune Response and Antioxidant Enzyme Activity in the Hepatopancreas of Prawns

The dietary intake of β -1,3-glucan significantly impacts the activities of immune and antioxidant-related enzymes in prawn hepatopancreas ($p < 0.05$; Figure 3). Feeding prawns with diets B1, B2, B3, and B4 resulted in a decrease in MDA content in hepatopancreas compared to those fed on diet B0 ($p < 0.05$; Figure 3A). Prawns fed on diets B1, B2, and B4 exhibited higher SOD activities than those fed on diet B0 ($p < 0.05$; Figure 3B). Additionally, prawns fed on diet B1 showed a higher SOD activity compared to other groups ($p < 0.05$), while prawns fed on diet B3 did not show a significant difference in SOD activity compared to the control group.

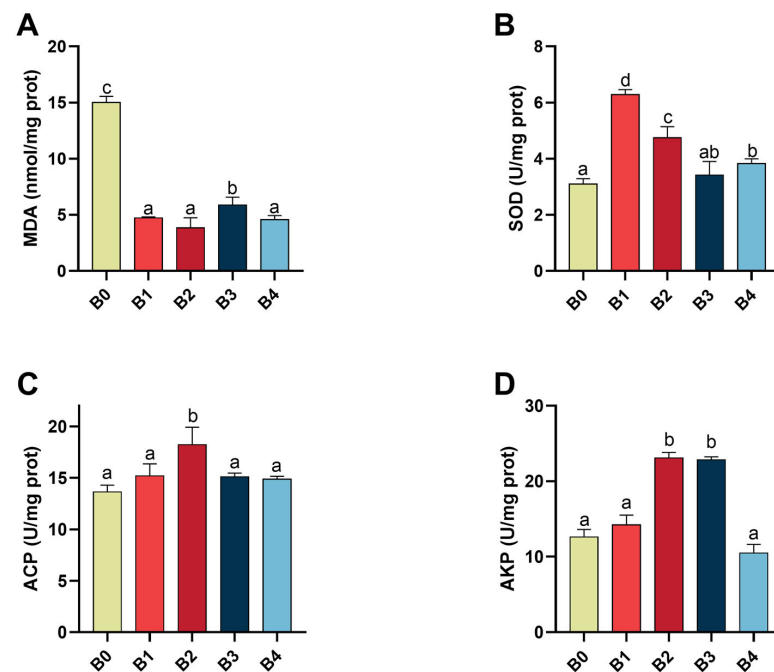


Figure 3. Antioxidant enzyme activities and innate immune response in the hepatopancreas of juvenile *M. nipponense* fed different concentrations of β -1,3-glucan. (A) Malondialdehyde (MDA) content, (B) superoxide dismutase (SOD) activity, (C) Acid phosphatase (ACP) activity, (D) Alkaline phosphatase (AKP) activity. Values are means for three replicates for each treatment \pm SD. Bars with different superscript letters are statistically significant ($p < 0.05$).

Prawns fed on diet B2 demonstrated higher ACP activity in the hepatopancreas than all other experimental diets ($p < 0.05$; Figure 3C). Furthermore, there was no significant difference in ACP activity among prawns fed on other diets. Prawns fed on both diet types B2 and B3 displayed an increase in AKP activity compared to all other experimental diets ($p < 0.05$; Figure 3D). Moreover, there was no statistically significant difference observed in the AKP activity of prawns that were administered either B2 or B3 diets.

3.4. The mRNA Expression of Genes Related to PRRs and Apoptosis

The mRNA expression of genes related to PRRs and apoptosis in the hepatopancreas was significantly affected by feeding prawns with different levels of dietary β -1,3-glucan ($p < 0.05$; Figure 4). Compared to the control diet, prawns fed on the β -1,3-glucan diet showed an upregulation in mRNA relative expression of the *LGBP* gene in the hepatopancreas ($p < 0.05$; Figure 4A). Furthermore, compared to those fed on all other diets, prawns fed on diet B2 exhibited an increase in mRNA expression of the *LGBP* gene ($p < 0.05$). The *lectin* gene expression in prawns fed on diets B1 and B2 was elevated compared to those fed on diets B0, B3, and B4 ($p < 0.05$; Figure 4B). Additionally, the *lectin* gene expression was increased in prawns fed on diet B2 compared to those fed on diet B1 ($p < 0.05$; Figure 4B).

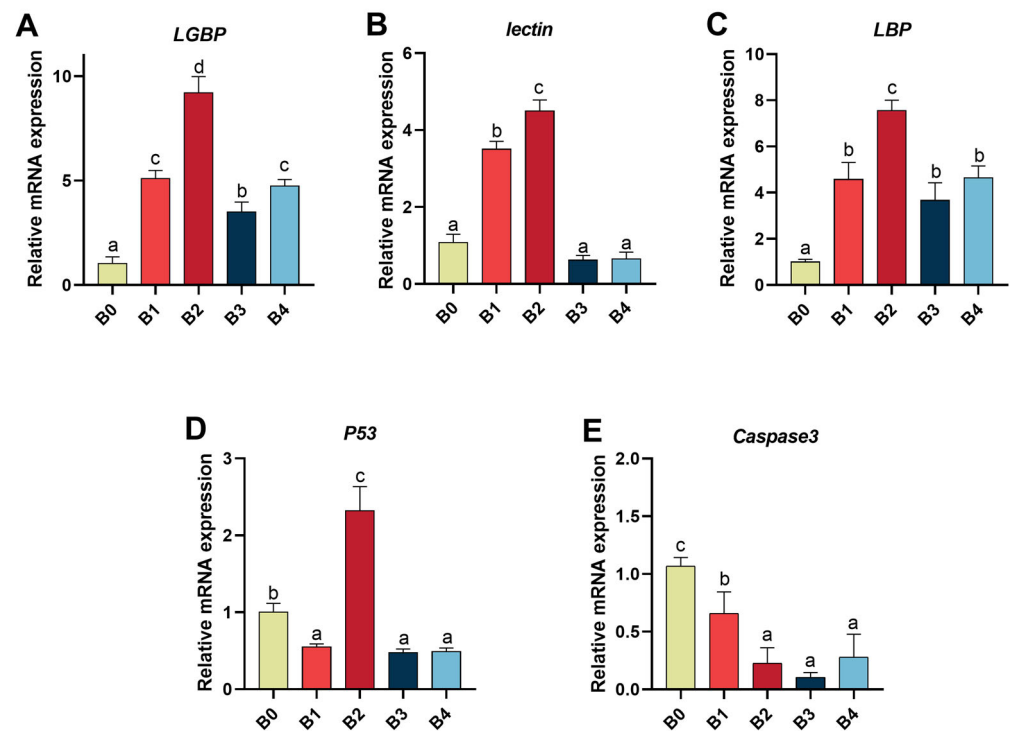


Figure 4. Effect of β -1,3-glucan on mRNA expression of immune-related genes in the hepatopancreas of juvenile oriental river prawns (*M. nipponense*) fed diets supplemented with different concentrations of β -1,3-glucan for eight weeks. (A) lipopolysaccharide and β -1,3-glucan binding protein (*LGBP*), (B) lectin, (C) lipopolysaccharide-binding protein (*LBP*), (D) tumor suppressor protein p53 (*p53*), (E) cysteine-aspartic acid protease 3 (*caspase 3*). Values are means \pm SD for three replicates for each treatment. Bars with different superscript letters are statistically significant ($p < 0.05$).

The prawns fed on the β -1,3-glucan diet upregulated the mRNA relative expression of the *LBP* gene in the hepatopancreas compared to the control diet ($p < 0.05$; Figure 4C). Moreover, prawns fed on diet B2 exhibited an enhancement in mRNA expression of the *LBP* gene compared to those fed on all other diets ($p < 0.05$).

Interestingly, *P53* gene mRNA expression was down-regulated in both diets B1, B3, and B4 compared to diet B0 ($p < 0.05$; Figure 4D). However, prawns fed on diet B2 showed an increase in mRNA expression of the *P53* gene compared to those fed on all other diets

($p < 0.05$). Prawns fed with the β -1,3-glucan diet exhibited an increase in *Caspase3* mRNA expression compared to those in the control group ($p < 0.05$; Figure 4E).

3.5. Gut Microbiota

The groups B0, B2, and B4 were selected for high-throughput sequencing analysis based on the increasing dietary β -1,3-glucan additions. The results presented in Table 4 revealed a total of 274,131 sequences across the three groups (B0, B2, and B4), with an average of 91,377 sequences per sample. The coverage within each group exceeded 99.8%, indicating that the obtained 16S rRNA gene sequences represented the majority of bacterial populations present in the samples. No significant difference was observed in sequence numbers among the B0, B2, and B4 groups. Similarly, there were no significant variations in Chao indices or Shannon and Simpson diversity measures between the three groups.

Table 4. Diversity of intestinal flora of juvenile oriental river prawn (*M. nipponense*) in groups B0, B2, and B4.

Diet	Sequences	Chao	Shannon	Simpson	Coverage (%)
B0	89,538 ± 16,475	925.66 ± 97.81	4.53 ± 0.12	0.045 ± 0.02	99.87 ± 0.06
B2	94,935 ± 27,033	763.73 ± 58.44	4.47 ± 0.21	0.033 ± 0.01	99.93 ± 0.02
B4	89,658 ± 56,480	829.08 ± 77.77	4.40 ± 0.12	0.044 ± 0.01	99.90 ± 0.03

The value is the average of the three replicates for each treatment ± SD.

Using a relative abundance threshold of 1%, we classified the dominant bacterial phyla and genera in the gut of prawns fed on experimental diets (Tables 5 and 6) based on phylum (Figure 5A) and genus (Figure 5B). The dominant phyla observed were Proteobacteria, followed by Cyanobacteria, Actinobacteria, and Patescibacteria (Table 5; Figure 5A). Although not statistically significant, a slight decrease in Actinobacteria and Patescibacteria abundance was observed in prawns fed diets B2 and B4 compared to those fed diet B0 (Table 5).

Table 5. The relative abundance of major phylum in intestinal flora of juvenile *M. nipponense* in groups B0, B2, and B4 fed experimental diets.

Phylum	Experimental Diet		
	B0	B2	B4
Proteobacteria	39.31 ± 14.65	56.99 ± 12.11	54.01 ± 13.98
Cyanobacteria	23.27 ± 7.37	15.55 ± 17.82	24.44 ± 13.09
Actinobacteria	10.24 ± 4.10	5.90 ± 2.56	5.98 ± 0.51
Patescibacteria	8.40 ± 4.05	5.67 ± 2.34	4.39 ± 1.06

The value is the average of the three replicates for each treatment ± SD.

Table 6. The relative abundance of major genera in intestinal flora of juvenile *M. nipponense* in groups B0, B2, and B4 fed experimental diets.

Genus	Experimental Diet		
	B0	B2	B4
<i>Chloroplast_norank</i>	12.03 ± 8.31	10.48 ± 12.76	10.60 ± 2.89
<i>Candidatus_Hepatincola_norank</i>	10.37 ± 10.87	7.29 ± 3.28	16.91 ± 2.87
<i>Rhodobacteraceae_unclassified</i>	2.24 ± 0.90 ^a	8.54 ± 3.78 ^b	4.20 ± 3.21 ^{ab}
<i>Saccharimonadales_norank</i>	7.44 ± 3.66	5.14 ± 2.07	4.00 ± 1.23
<i>Rhodobacter</i>	2.48 ± 0.67 ^a	9.67 ± 3.18 ^b	6.36 ± 4.47 ^{ab}

The value is the average of the three replicates for each treatment ± SD. The presence of different superscript letters in the same column indicates significant differences ($p < 0.05$).

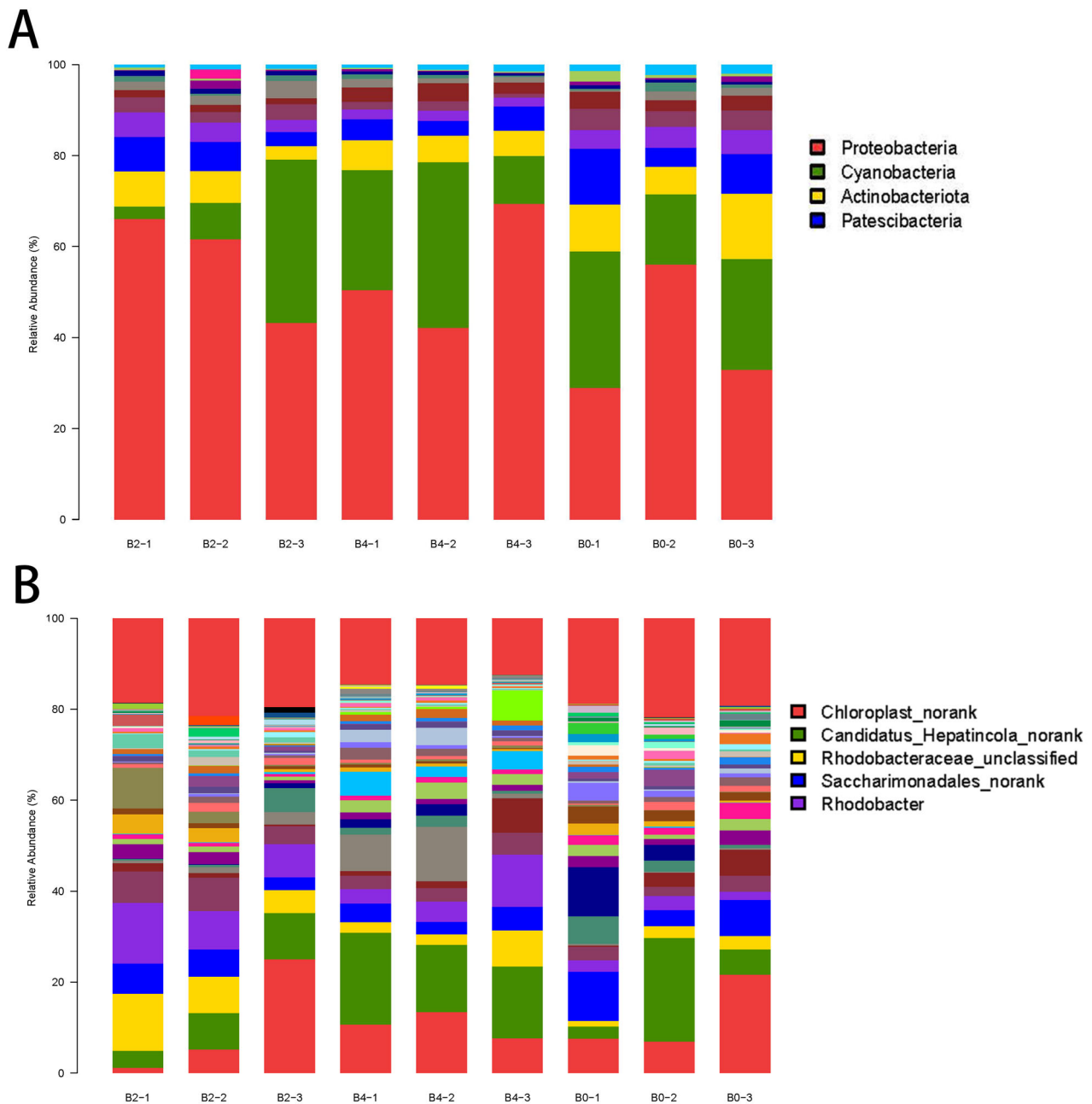


Figure 5. Histogram of intestinal microorganism distribution in juvenile oriental river prawns (*M. nipponense*) supplemented with different dietary concentrations of β -1,3-glucan. (A) phylum taxonomic level; (B) genus taxonomic level.

The dominant genera in the gut of the prawns fed on the experimental diets were primarily *Chloroplast_norank*, followed by *Candidatus_Hepatincola_norank*, *Rhodobacteraceae_unclassified*, *Saccharimonadales_norank*, and *Rhodobacter* (Table 6; Figure 5B). There were no significant differences observed in the abundance of *Chloroplast*, *Candidatus*, and *Saccharimonadales_norank* among prawns fed on all diets. Interestingly, the prawns fed on diet B2 showed a higher abundance of *Rhodobacteraceae_unclassified* and *Rhodobacter* compared to those fed on diets B0 and B4 ($p < 0.05$).

4. Discussion

In the present study, the inclusion of dietary β -1,3-glucan enhanced the weight gain and survival rate of *M. nipponense*. The improved growth performance and survival rate

observed in *M. nipponense* may be attributed to enhanced absorption and accumulation of nutrients as well as improved innate immune response facilitated by β -1,3-glucan supplementation. Despite being underutilized in practical agriculture due to a lack of scientific strategies for its application, our study demonstrates that incorporating β -1,3-glucan into feed rations can effectively enhance growth performance and survival rates, highlighting its potential for superior production applications. Similar findings have been reported with *L. vannamei* where the addition of 0.02% or 0.04% β -glucan resulted in increased weight gain [35,54]. Furthermore, β -glucan has also shown significant improvements in weight gain and survival rate of *Penaeus monodon* [55]. Notably, the present study revealed a trend wherein the survival rate of *M. nipponense* initially increased but then decreased with increasing levels of added β -1,3-glucan. Therefore, we recommend 0.18% β -1,3-glucan in *M. nipponense* diets to ensure optimal survival rate.

Oxygen is known to produce ROS, which are harmful to living organisms. [56]. Lipid peroxidation is a common injury phenomenon induced by ROS, resulting from the oxidation of polyunsaturated fatty acids that are susceptible to attack by free radicals [57]. Therefore, organisms possess enzymatic antioxidant defenses to minimize the detrimental effects of ROS [58]. SOD plays a crucial role as an important component of the antioxidant enzyme system in biological systems and acts as a key enzyme in the elimination of ROS [59]. MDA serves as an end product of lipid peroxidation and indirectly reflects the tissue damage caused by peroxidation [60]. Hence, MDA content and SOD activity in the hepatopancreas can serve as an indicator for assessing the antioxidant capacity of prawns. In this study, feeding prawns with a β -1,3-glucan diet resulted in increased SOD activity but decreased MDA levels in *M. nipponense*. These findings demonstrate that β -1,3-glucan enhances the antioxidant capacity of *M. nipponense*. Similar results regarding enhanced antioxidant capacity after β -1,3-glucan supplementation have also been reported in swimming crabs (*Portunus trituberculatus*) [61] and *Apostichopus japonicus* [62]. Additionally, our experiment results indicate that β -1,3-glucan can increase AKP and ACP activities in the hepatopancreas of *M. nipponense*. The activity level initially increases with increasing amounts of added β -1,3-glucan but eventually decreases thereafter; with 0.2% being associated with the highest observed activity level among all tested concentrations. This observation can be attributed to crustaceans possessing a large and complex innate immune system [63], wherein AKP and ACP play vital roles as phosphatases involved in non-specific immunity responses [64]. Previous studies have shown increased ACP and AKP activities when red swamp crayfish (*Procambarus clarkii*) [36] and Ussuri catfish (*Pseudobagrus ussuriensis*) [65] were fed diets supplemented with β -glucan. These findings demonstrate that the inclusion of appropriate β -1,3-glucan enhances both antioxidant capacity and nonspecific immunity in *M. nipponense*. Notably, the improved non-specific immunity holds promise for reducing antibiotic usage during infectious disease outbreaks in prawn farming, thereby promoting sustainable development of prawn aquaculture.

Cumulative survival following bacterial infection serves as a straightforward indicator to assess disease resistance in aquatic animals and evaluate the efficacy of cultivating biological health and enhancing immunity [12]. Importantly, *M. nipponense* fed with β -1,3-glucan diets exhibited a distinct increase in cumulative survival after *A. hydrophila* infection; moreover, cumulative survival displayed an increasing trend followed by a decrease upon the addition of β -1,3-glucan. These observations suggest that appropriate amounts of addition of β -1,3-glucan can enhance the disease resistance of *M. nipponense*. Similarly, *Micropterus salmoides* showed higher survival after *Aeromonas schubertii* infection when fed diets supplemented with β -glucan [66], while juvenile tiger shrimp demonstrated increased survivability after WSSV infection when fed diets enriched with β -glucan [67]. Collectively, these results indicated that to some extent β -1,3-glucan has the potential to mitigate aquatic disease outbreaks caused by pathogenic microorganisms. The fundamental role of the immune system is to distinguish between “self” and “non-self”. Crustaceans, known for their innate immunity, heavily rely on the recognition of “non-self” [25]. Recent findings indicate that this recognition is facilitated by certain PRRs, which can be soluble or

membrane-bound [68]. A critical step in initiating the innate immune response involves the mutual recognition and interaction between PRRs and PAMPs present on the surface of pathogens and absent in the host [69]. β -1,3-glucan found on fungal cell walls represents an important class of PAMPs [70], capable of binding to common crustaceans PRRs such as *LBP* [11], *bGBP* [14], *LGBP* [12], and *lectin* [71]; this interaction subsequently triggers proPO (Prophenoloxidase), a widely distributed enzyme in blood and inner tissues of carapace, leading to activation of proPO-AS [41]. Therefore, quantitatively analyzing mRNA expression levels of *LBP*, *LGBP* and *lectin* in prawn hepatopancreas can serve as an effective tool for assessing prawn immune response status. Our results further demonstrate that feeding the prawns with β -1,3-glucan upregulates mRNA expression levels of *LBP*, *LGBP*, and *lectin* genes. Interestingly, numerous studies have investigated the relationship between β -glucan and aquatic animals, with PO activity often being associated with certain PRRs-related genes. Therefore, we hypothesized that upregulating the expression of *LBP*, *LGBP*, and *lectin* genes could enhance the immunity of *M. nipponense* by increasing PO activity. For instance, *L. vannamei* fed a diet supplemented with β -glucan exhibited simultaneous enhancements in both PO activity and *LGBP* gene expression [12], while *Macrobrachium rosenbergii* fed a similar diet showed concurrent increases in both PO activity and *lectin* gene expression [72]. Furthermore, our study found that the β -1,3-glucan group had a significantly higher cumulative survival rate 72 h after *A. hydrophila* infection compared to the control group. Hence, we postulated that *M. nipponense* could activate proPO-AS by enhancing the expression of *LBP*, *LGBP*, and *lectin* genes through β -1,3-glucan supplementation to resist *A. hydrophila* infestation. Previous research has also demonstrated significant induction of crustacean *LGBP* transcripts following challenges with bacteria such as *Vibrio harveyi* [73], *Vibrio alginolyticus* [74], and *Vibrio anguillarum* [75]. According to the results of the present study, the addition of moderate amounts of β -1,3-glucan to the diet improved the non-specific immunity of *M. nipponense*.

Caspases are closely associated with eukaryotic apoptosis and play a role in regulating cell growth, differentiation, and apoptosis [76]. Apoptosis is considered to be a form of programmed cell death that primarily functions to eliminate damaged, dangerous, and non-functional cells in order to maintain organismal health [77,78]. The caspase protease family consists of numerous members, among which downstream *caspase3* acts as the executors of apoptosis by enzymatically cleaving specific proteins [79]. In this study, the gene expression of *M. nipponense caspsae3* was significantly reduced in the group fed β -1,3-glucan compared to the control group. Therefore, it can be inferred that the addition of β -1,3-glucan to feed can attenuate apoptosis. *P53* is a protein involved in inhibiting cell cycle progression in DNA-damaged cells [80], which is regulated by activating cellular repair mechanisms for DNA damage repair or inducing apoptosis if repair is not feasible. It plays a pivotal role in gene expression related to cell cycle, genetic stability maintenance, and apoptosis [81]. In our experiment, the inclusion of 0.05%, 0.2%, and 0.4% β -1,3-glucan significantly downregulated *P53* gene expression in prawns consistently with *caspase3* trend. We speculate that the addition of β -1,3-glucan to the diet could attenuate apoptosis in *M. nipponense* by enhancing the nonspecific immunity characteristic of crustaceans, which is consistent with the findings shown above for ACP and AKP. Furthermore, a correlation between the enhancement of specific immunity and the reduction in apoptosis, as indicated by ACP and AKP levels, has been observed in several other studies [82,83]. Interestingly, the inclusion of 0.1% β -1,3-glucan significantly up-regulated *P53* gene expression possibly due to its upstream position relative to *caspase3*, as well as its involvement in antioxidant defense regulation, DNA repair processes, and apoptotic pathways [84].

The intestine is an important organ for the digestion and absorption of nutrients in shrimp [85,86]. Gut microorganisms are essential for nutritional digestion and absorption, as well as immune function and disease resistance [87]. Feeding probiotics to *Oreochromis niloticus* has been shown to enhance intestinal villi development, growth performance, and mucosal immunity [88]. Similarly, improving gut flora has been found to enhance immune responses and disease resistance in various finfish species [89]. Identifying up- and down-

regulated microorganisms is an important step in understanding how β -glucan affects the gut microbiota and how this change in the microbiota affects the host by altering nutrients, the immune system, and compounds absorbed through the epithelial barrier [90]. In this study, the addition of β -1,3-glucan did not affect the gut flora abundance at phylum level, we speculate that the addition of β -1,3-glucan at the phylum level maintains the stability and diversity of the intestinal flora of *M. nipponense*. At the genus level, this experiment demonstrated that the addition of β -1,3-glucan significantly increased the abundance of *Rhodobacteraceae_unclassified* in the intestinal tract of *M. nipponense*. Consistent with previous studies on β -glucan, it has been consistently observed that supplementation with β -glucan enhances the abundance of *Rhodobacteraceae*, which may represent a key characteristic associated with competitive advantage induced by β -glucan [91,92]. Moreover, a prior investigation on *N. californicus* revealed that *Rhodobacteraceae* can serve as a probiotic in the diet of *N. californicus* and improve resistance to acute low-salinity challenge in shrimp *Penaeus vannamei* through higher survival rates and elevated levels of T-AOC activity as well as SOD, HSP70, and Relish gene transcripts [93]. Therefore, dietary supplementation with β -1,3-glucan could enhance *Rhodobacteraceae* abundance for gut health improvement in *M. nipponense*. However, further exploration is required to fully understand this potential mutualistic relationship between *M. nipponense* and specific taxa within its intestinal flora.

5. Conclusions

The present study demonstrated that the addition of appropriate amount of β -1,3-glucan to the diet improved the growth performance, survival, antioxidant capacity, and intestinal health of *M. nipponense*, within the concentration range of β -1,3-glucan set in this experiment. Moreover, β -1,3-glucan activation of pattern recognition receptors initiated diverse immune responses to effectively enhance the survival and disease resistance in *M. nipponense*.

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