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Posted Date: 29 September 2023

doi: 10.20944/preprints202309.2037.v1

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Article

First Investigation of the Optimal Vaccination Timing for Nile Tilapia (*Oreochromis niloticus*) Larvae against *Streptococcus agalactiae*

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Abstract: To investigate the early immune responses and explore the optimal vaccination periods, Nile tilapia at 1, 7, 14, 21, 28, 35, and 42 days after yolk sac collapse (DAYC), were immersed in formalin-killed *Streptococcus agalactiae* vaccine (FKV-SA). The results found that specific IgM by ELISA in 21 DAYC (0.108 g) was first detected at 336 h after vaccination (hav), whereas 28–42 DAYC (0.330–0.580 g), could be initially detected at 24 hav. qRT-PCR analysis of the *TCRβ*, *CD4*, *MHCIIα*, *IgHM*, *IgHT*, and *IgHD* genes at 21–42 DAYC immunized with FKV-SA immersion route for 24, 168 and 336 hav, revealed that most immune-related genes were significantly higher in the vaccinated larvae in all DAYCs than the control larvae ($P < 0.05$) after 336 hav. Immunohistochemistry demonstrated the stronger signals of IgM in the gills, head kidney, and intestine tissues at 21, 28, and 35 DAYC were observed in all vaccinated larvae compared with the control. Interestingly, at all DAYCs, FKV-SA larvae exhibited significantly higher survival rates and an increased relative percent survival (RPS) than the control after challenge with viable *S. agalactiae*, particularly in larval fish that were immunized with FKV-SA for 168 and 336 hav ($P < 0.05$).

Keywords: Nile tilapia; *Oreochromis niloticus*; *Streptococcus agalactiae*; formalin-killed vaccine; immunocompetence

1. Introduction

Oreochromis niloticus is classified as a group of vertebrates and is a species of freshwater cichlid known as Nile tilapia. Like many countries worldwide, it is an economically important fish with the highest value and productivity in Thailand's freshwater aquaculture [1]. Due to the increasing demand for aquatic protein as a food source, the production capacity per unit area has increased with intensive aquaculture to provide more food in both quantity and quality. This results in a large amount of leftover feed and excreta, causing the accumulation of organic and inorganic substances in the culture pond, which always causes sudden changes in culture pond conditions, especially chemical, biological, and physical factors of water quality. These fluctuations always induce fish to be stressed, weakened, and more susceptible to various pathogens. Among these diseases, the

pathogenic bacteria *Streptococcus* spp., especially *S. agalactiae* and *S. iniae*, have caused streptococcosis diseases that severely damage Thailand's tilapia aquaculture industry, similar to other countries in temperate and tropical regions [1-4].

All bacteria in the genus *Streptococcus* belong to Streptococcaceae. These gram-positive bacteria are classified as lactic acid bacteria, are pathogenic and can cause high mortality rates of up to 75% with economic impact loss in tilapia cultures and other fish species [5,6]. Generally, antibiotics and chemicals are used to control bacterial diseases. However, the application of these agents causes several consequent effects, such as contamination in the environment and fish products, that further affect beneficial microorganisms and health problems in consumers. Additionally, drug resistance of pathogenic bacteria is always observed and eventually causes failures of disease control of these pathogens in fish culture systems [7,8].

To overcome these disease concerns, vaccines are used to provide an effective strategy for disease control. The development of vaccines in fish aquaculture has now flourished, and many commercial vaccines have been launched in several of the most economically important fish species at commercial scales [9-11], including Nile tilapia [12-14]. Critical problems with this strategy remain and are not practical, especially for the vaccination route. Among the vaccine administrations (injection, oral route, and immersion), injection is the most effective based on immune responses and protective efficacy [15,16], while oral and immersion vaccinations are more practical but less effective than injection administration. Therefore, oral and immersion vaccinations are the major practices for increasing vaccine efficacy, especially immersion vaccination. Compared with other methods, the immersion route is required since it has many advantages, including being less labor intensive, minimizing stress, and being administered in various fish sizes, particularly small-size fish with less vaccine quantity during application [17,18].

Based on current information, immersion vaccination in the early stages of fish is not so successful. One of the biggest problems is understanding the optimal stages that duly prompt exploration of the maximal immune responses, as little is known, especially in Nile tilapia. Therefore, we aimed to investigate the early immune responses of Nile tilapia larvae following immersion vaccination with inactivated *S. agalactiae*. Our investigation encompasses an assessment of specific IgM antibody responses, an analysis of immune gene expression, and an evaluation of vaccine efficacy through challenge tests. The basic knowledge arising from this study will be an important part of creating an effective strategy to prevent severe losses caused by various disease outbreaks in Nile tilapia. These results are also essential for the Nile tilapia aquaculture industry, especially for enhancing production and reducing the cost and negative side effects from antibiotic and chemical applications, which are crucial to sustainable Nile tilapia aquaculture.

2. Materials and Methods

2.1. Experimental fish and husbandry

A total of 35 male, apparently healthy (tested negative for pathogens) Nile tilapia (*Oreochromis niloticus*) approximately 6 months of age, weighing 200 ± 25 g and with a length of 20.6 ± 1.5 cm and 96 females weighing 200 ± 25 g and with a length of 18.8 ± 1.3 cm were bred in a 5×10 -m² cement tank containing freshwater with a full aeration system at the Fisheries Research Station, Kamphaengsaen campus, Nakornprathom province, Thailand. During spawning, tilapia eggs were collected from female fish and kept in the flow-through hatchery system until they reached the fifth stage [yolk sac collapse (YSC) larvae]. Then, 7,000 larvae at the same stage were further moved to the larval nursery system. Fish larvae were acclimatized in a 3,000-L tank with aerated freshwater at a temperature of 28 ± 2 °C, dissolved oxygen (DO) 5 ± 1.5 mg/L, and pH 7.8 ± 1.0 . Water quality was monitored and exchanged on alternate days. The fish were fed commercial larval feed (Hi-Grade 9006T, Charoen Pokphand, Thailand) with an approximate crude protein content of 42% at 10% of the body weight 3 times a day for 42 days (42 DAYC).

The experimental procedures and the use of animals were approved according to the guidelines of the Animal Welfare Committee of Kasetsart University (ACKU65-FIS-016).

2.2. Bacterial cultivation and preparation of formalin-killed *S. agalactiae* vaccine (FKV-SA)

2.2.1. Bacterial preparation

Streptococcus agalactiae bacterium serotype III was previously isolated from diseased Nile tilapia and identified by the Center of Excellence in Aquatic Animal Health Management, Department of Aquaculture, Faculty of Fisheries, Kasetsart University, Bangkok, Thailand. A single colony of the target strain was inoculated into tryptic soy broth (TSB, Becton, Dickinson and Company, USA) and incubated overnight at 37 °C for 24 h with shaking at 120 rpm. Pellets were obtained by centrifugation at 5,000 × g for 12 min and washed twice with 0.85% NaCl, and then the supernatant was discarded. Bacterial concentrations were monitored for target absorbance using a spectrophotometer (Milton Roy, USA) at an optical density of 600 nm to reach 1.0 to obtain a final bacterial cell concentration of approximately 1×10^9 CFU/ml. This concentration was further diluted for use in the other experiments, and the number of bacteria was always confirmed by spreading on tryptic soy agar (TSA).

2.2.2. Formalin-killed *S. agalactiae* vaccine preparation (FKV-SA)

The above-prepared bacterial cells were inactivated in 1.0% (v/v) formalin solution and incubated overnight at 4 °C. Bacterial inactivation was verified by spreading 100 µL on TSA (Becton, Dickinson and Company, USA) and incubating at 37 °C for 16-24 h. When no colonies were observed, the inactivated bacterial suspension was prepared as described above, and the bacterial pellet was washed twice with 0.85% NaCl and centrifuged for 12 min at 5,000 g and 4 °C for 12 min. Bacterial cell density was measured by previous methods, and a final bacterial cell concentration of approximately 1×10^9 CFU/ml was maintained at 4 °C until use.

2.3. Efficacy of the *S. agalactiae* immersion vaccine (FKV-SA) in Nile tilapia larvae

2.3.1. Experimental design

According to the factorial experiment in a completely randomized design (CRD), a 2×5 factorial experiment with two replicates was designed with two factors (vaccination and days after vaccination). Seven different Nile tilapia larvae at 1, 7, 14, 21, 28, 35, and 42 DAYC in section 2.1 were used in this study. Nile tilapia larvae were randomly divided from storage tanks into four 500-liter L tanks, each containing 500 fish. Tanks 1 and 2 were assigned as control groups 1 and 2, and tanks 3 and 4 were assigned as vaccinated groups 1 and 2 (FKV-SA1 and 2). A total of 28 experimental tanks were set up. The fish at different larval stages were weighed and scaled for growth parameter determination.

2.3.2. Immersion vaccination

The vaccine prepared as described in 2.2.2 was diluted to a concentration of 1×10^7 CFU/mL. At each fish larval stage, 4 20-L tanks were prepared. Tanks 1 and 2 were the control group immersed in 0.85% NaCl, and tanks 3 and 4 were immersed in formalin-killed *S. agalactiae* vaccine (FKV-SA) at a concentration of 1×10^7 CFU/mL for 2 h. All tanks were continuously aerated during immunization, and water temperatures were maintained at 28 ± 2 °C. After vaccination, fish were returned to their nursery tanks.

2.3.3. Sample collection

The whole bodies of 18 tilapia larvae in the vaccinated and unvaccinated groups (9 fish per tank) were randomly collected at 1, 7, 14, 21, 28, 35, and 42 DAYC at 0, 6, 24, 168, and 336 h after vaccination (hav). Six fish were kept in TRIzol™ reagent (Thermo Fisher Scientific, USA) and kept at -80 °C for gene expression studies for a short period, and the other 6 whole fish samples were kept at -20 °C for assessment of serum antibody by ELISA. The 6 remaining whole tilapia larvae were preserved in 10% buffer formalin solution at room temperature for immunohistochemical analysis.

2.3.4. Measurement of the fish weight and size prior to and post-vaccination

Two growth parameters of each experimental group were evaluated: body weight and total length. Six fish from each control and treatment group were randomly selected in each DAYC at 1, 7, 14, 21, 28, 35, and 42 and at 0, 6, 24, 168, and 336 h after inoculation to measure weight and length.

2.3.5. Assessment of specific IgM levels to FKV-SA by ELISA

Fish larvae from the previous section were extracted for protein solution using previously described methods [19]. The ELISA microplates were first coated with carbonate-bicarbonate coating buffer, pH 9.6, for 1 h, and the supernatant was discarded and washed three times with 1× Low Salt Wash Buffer (LSWB). One hundred microliters of the FKV-SA at 1×10^9 CFU/mL in PBS was added to the wells of the plates. The plates were incubated overnight at 4 °C, and 50 µL of 0.05% glutaraldehyde was added for 20 min at room temperature (RT). Then, the glutaraldehyde was discarded, and the cells were washed three times with 1× LSBW. The microplate wells were further blocked with 100 µL of 1% bovine serum albumin (BSA) blocking buffer (Sigma A9647) for 2 h at RT and then washed five times with 1× high salt wash buffer (HSWB) before adding 100 µL of the sampled fish supernatant to the microplate wells and incubating overnight at 4 °C. After washing the plate five times with HSWB, anti-tilapia IgM monoclonal antibody (1:200 v/v) (Marine Leader, Thailand) was added for 2 h. The plates were washed five times with HSWB and incubated for 5 min on the last wash, and 100 µL of conjugated anti-mouse IgG-HRP (Sigma Aldrich, USA) diluted at 1:3000 v/v was added and incubated for 1 h at RT. The plates were washed five times and incubated for 5 min on the last wash. One hundred microliters of TMB (3',3',5',5' – tetramethylbenzidine dihydrochloride) was added to each well, and the plates were incubated at RT for 10 min in the dark. The reaction was stopped with 100 µL of H₂SO₄ (Sigma–Aldrich 320501). The absorbance of the reaction was measured with an iMark™ Microplate Absorbance Reader (Bio-Rad iMark™ microplate reader). The serum antibody levels specific to the FKV-SA antigen of fish larvae at each DAYC and the day after vaccination were determined using the absorbance values at 450 nm.

2.3.6. Determination of immune-related gene expression of tilapia larvae after immersion vaccination

Approximately 40 mg of visceral organs of six fish from each treatment were randomly sampled from each replicate at 21, 28, 35, and 42 days at 24, 168, and 336 h after postimmersion vaccination. Tissues were placed in 1.0 mL of TRIzol™ reagent (Thermo Fisher Scientific, USA) according to the manufacturer's instructions. Total RNA was isolated from the tissues, and the RNA concentration and purity were determined using a Nanodrop spectrophotometer (Bio-Rad Laboratories, Hercules, California, USA). One microliter of 1000 ng/µL total RNA was then used as a template to synthesize first-strand cDNA using the ReverTra Ace® qPCR RT Master Mix with gDNA Remover Kit (TOYOBO, Japan). The first strand cDNA synthesis product was stored at -80 °C for further analysis.

Immune-related gene expression for *TCRβ*, *CD4*, *MHCIIα*, *IgM*, *IgT*, and *IgD* was followed using the experiment by Bunnoy et al. (2023) [19]. The specific primers are shown in Table 1. The qRT-PCR assay was performed using Brilliant III Ultra-Fast SYBR® Green (Agilent, CA, USA) in an AriaMx Real-Time PCR system (MY18135206). qPCR was performed in a total volume of 20 µL containing 10 µL of 2 × SYBR Green QPCR Master Mix, 2 µL of 0.5 mM forward and reverse primers, and 1 µL of cDNA template, adjusting the reaction mixture to a final volume of 20 µL with distilled water, and the thermal cycler program was run at 95 °C for 5 min; 40 cycles of 95 °C for 30 s, 55 °C for 30 s, and 72 °C for 90 s; and a final extension at 72 °C for 10 min as previously described [20]. All reactions were performed in triplicate. The expression levels of each sample were described by the relative fold change to that of *β-actin* using the $2^{-\Delta\Delta CT}$ method according to the protocol by Livak and Schmittgen (2001) [21].

Table 1. Primers were used in this study to determine gene expression levels in Nile tilapia (*Oreochromis niloticus*).

Genes	Primer names	Nucleotide sequences (5'→3')	Amplicon size (bp)	Temperature (°C)	References
β-actin	On_β-actin	F: ACAGGATGCAGAAGGAGATC ACAG R: GTACTCCTGCTTGCTGATCCACAT	155	60	[19]
T cell receptor β-chain (constant region)	On_TCR β	F: GGACCTTCAGAACATGAGTGCAG R: TCTTCACGCGCAGCTTCATCTGT	164	60	Nithikulwong <i>et al.</i> (2012) [75]
Cluster of differentiation 4 (CD4)	On_CD4	F: GCTCCAGTGTGACGTGAAAR: TACAGGTTTGAGTTGAGCTG	150	60	[19]
Major histocompatibility complex (MHC) class IIα molecules	On_MH C-IIα	F: CAGTGTTTGATGTGTTTTTCAGR: CTCTTCACCATCCAGTCCA	100	60	[19]
Immunoglobulin M heavy chain (<i>IgHM</i>)	On_IgM	F: GGATGACGAGGAAGCAGACTR: CATCATCCCTTTGCCACTGG	122	60	[19]
Immunoglobulin T heavy chain (<i>IgHT</i>)	On_IgT	F: TGACCAGAAATGGCGAAGTCTG R: GTTATAGTCACATTCTTTAGATTACC	136	60	[19]
Immunoglobulin D heavy chain (<i>IgHD</i>)	On_IgD	F: AACACCACCCTGTCCCTGAAT R: GGGTGAAAACACATTCCAAAC	127	60	Wang <i>et al.</i> (2016) [76]

2.3.7. Immunohistochemical analysis

Whole-body samples were randomly collected from six fish from each replicate tilapia larvae at days 21, 28, and 35 and at 0, 6, 24, 168, and 336 h after vaccination for immunohistochemical analyses.

The whole fish samples were preserved in a 10% buffer formalin fixative solution for 24 h. They were then dehydrated in a graded ethanol series and xylene before embedding in paraffin. Sections were mounted on glass slides and dried for 24 h at 37 °C. Slides were deparaffinized, rehydrated in xylene for 5 min (twice), absolute ethanol for 5 min (twice), 80% EtOH for 5 min, and 70% ethanol for 5 min, and the slides were washed with distilled water for 5 min. Then, the slides were pretreated for 30 min with 3% H₂O₂ in 70% ethanol in the dark at RT and then washed with PBS. The slides were incubated with 1% glycine for 30 min at RT. Antigens were retrieved by breaking the crosslinks between antigens and fixation or paraffin, boiling in citrate buffer, and sliding for 10 min on a hot plate and natural cooling at RT. Slides were washed with 1× PBST for 5 min (twice) and permeabilization by 0.4% Triton X-100 in 1× PBS for 5 min (twice). Slides were further blocked with 5% BSA (in 1X PBS) for 30 min in a humid box at RT and washed with 1× PBST for 5 min (twice). Slides were incubated at 37 °C for 3 h with primary antibody (IgM) diluted in PBS (1:100). After washing, the slides were incubated with GAM-HRP (secondary antibody) 1:1000 of 1×PBS and incubated at 37 °C for 2 h with 1× PBST for 5 min (3 times). Then, the bound antigen was visualized by incubating slides in DAB staining solution in the dark for 15 min. Slides were counterstained with Meyer hematoxylin and mounted in Permount™ mounting medium (PMM). Section images were collected using a compound light microscope (Olympus, MA, USA). DAB showing positive signals was stained brown, and nuclei were stained blue.

2.4. Survival rate and relative percent survival (RPS) of vaccinated Nile tilapia challenged with *S. agalactiae*.

2.4.1. Assessment of the time-dependent toxicity of the lethal concentration (LC₅₀) of *S. agalactiae* in Nile tilapia

Fish prepared as described in section 2.1 were used. The fish were divided into 6 experimental groups with 3 replicates (10 fish/replicates) as follows: group 1, which was the control group, was inoculated with PBS, and groups 2, 3, 4, 5, and 6 were inoculated with viable *S. agalactiae* at concentrations of 1×10⁴, 1×10⁵, 1×10⁶, 1×10⁷ and 1×10⁸ CFU/mL, respectively. The fish were exposed to the pathogens by immersion for 120 min and transferred to their tanks. The mortality rate was observed and recorded daily. The survival rate was then calculated for the median lethal concentration (LC₅₀) according to the methods by Finney (1971) [22] and Litchfield and Wilcoxon (1949) [23].

2.4.2. Challenge test of vaccinated Nile tilapia larvae

Fish were prepared as described in section 2.1. The experiment was divided into 4 experimental groups according to the tilapia larvae age: Groups 1-4 were days 21, 28, 35, and 42, respectively. Fifty larvae were maintained in 250 L glass tanks, with 3 repetitions.

Nile tilapia larvae were collected in each series of experiments as planned in section 2.4.1, with groups 1, 2, and 3 immersed in PBS pH 7.4 for 2 h and groups 4, 5, and 6 immersed in FKV-SA at a concentration of 1 × 10⁷ CFU/ml for 2 h. The larvae were then transferred to a 250 L glass. The experimental fish were then immersed in an *S. agalactiae* bacterial solution at the LC₅₀ value from section 2.4.1, which was 3.34 × 10⁵ CFU/mL, for 2 h at 24, 168, and 336 h after immunization. Fish mortality was recorded daily for 8 days after immunization. The survival rate and relative percent survival (RPS) were calculated as follows:

$$\text{Survival rate (\%)} = (\text{Number of fish remaining} / \text{Total number of fish}) \times 100$$

$$\text{RPS} = 1 - (\text{Mortality rate of vaccinated fish} / \text{Mortality rate of control fish}) \times 100.$$

2.5. Statistical and data analysis

IgM levels, immune-related gene expression, mortality, and growth performance were analyzed by two-way analysis of variance (ANOVA) and Duncan's multiple comparisons test to determine differences between groups. The results are expressed as the means ± SDs. The interactions between the determined factors were further analyzed. All statistical analyses were performed using IBM SPSS Statistics version 25 and GraphPad Prism 8.0.2. Survival analysis of Nile tilapia larvae at different

ages challenged with *S. agalactiae* was calculated using the Kaplan–Meier method. The level of statistical significance between the control and experimental groups in all parameters was indicated as $P < 0.05$.

3. Results

3.1. Levels of serum IgM specific to *S. agalactiae* by ELISA

The results of the inactivated *S. agalactiae* vaccine (FKV-SA) by immersion at time intervals of 0, 6, 24, 168-, and 336 hav showed no significant differences in the measurement of specific IgM levels against *S. agalactiae* at 1, 7, and 14 DAYC compared with the nonvaccinated control group ($P > 0.05$) (Figure 1A-C). However, in Nile tilapia larvae immunized with FKV-SA at 21 DAYC, a statistically significant increase in the IgM levels was observed at 336 hav compared with the nonvaccinated control group ($P < 0.05$) (Figure 1D). In addition, a significant increase in IgM levels specific to *S. agalactiae* was observed in Nile tilapia larvae at 28, 35, and 42 DAYC after vaccine immunization, especially at 24, 168, and 336 hav, respectively, compared with the nonvaccinated control group ($P < 0.05$) (Figure 1E-G). Furthermore, IgM levels increased in both vaccinated and unvaccinated treatments during every experimental period when overall vaccination periods were determined. However, no interaction between periods and vaccination treatments was observed throughout vaccination periods ($P > 0.05$).

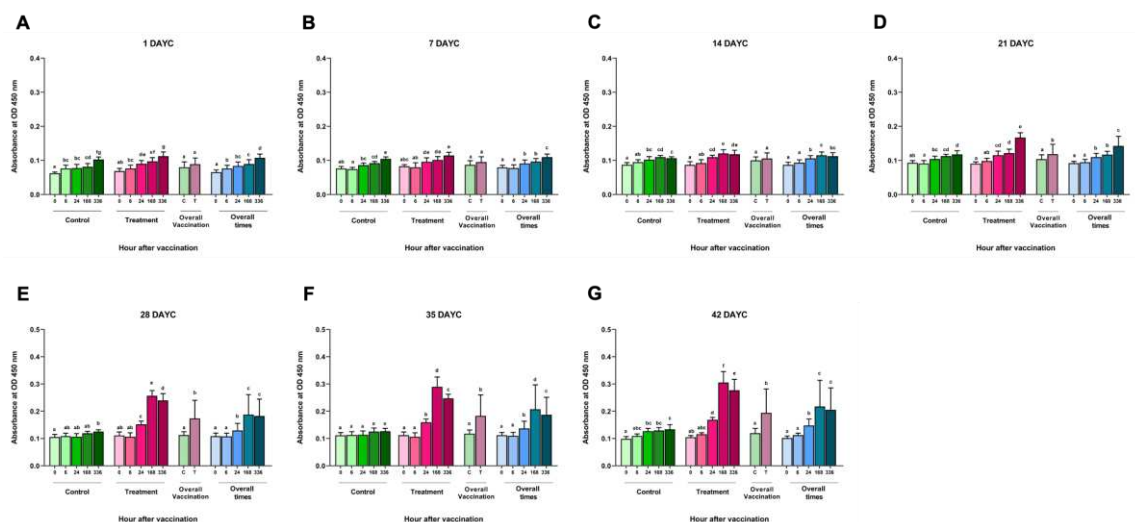


Figure 1. Specific IgM levels of FKV-SA Nile tilapia specific to *S. agalactiae*. The IgM levels were determined by ELISA at 1(A), 7(B), 14(C), 21(D), 28(E), 35(F) and 42(G) DAYC. Different letters on each bar denote significant differences ($P < 0.05$).

3.2. Immune-related gene expression

Based on the appearance of IgM levels in section 3.1, the relative gene expression of the immune-related genes *IgM*, *IgT*, *IgD*, *MHCII α* , *TCR β* and *CD4* in the whole body of Nile tilapia larvae was conducted at 21, 28, 35, and 42 DAYC and at 24, 168 and 336 hav. The relative expression of the *TCR β* gene in fish larvae at 21 DAYC initially demonstrated differences between the vaccinated and nonvaccinated groups at 336 hav. However, fish larvae at 28-42 DAYC and immunized with FKV-SA were significantly found to express the *TCR β* gene at higher levels than control fish (Figure 2A-2D). This agreed with the overall vaccination, in which the expression of the *TCR β* gene was first found to be significantly different at 28 DAYC larvae. When the effects of overall vaccination times were considered, significantly higher levels of the *TCR β* gene from 28-42 DAYC larvae at 168 and 336 hav were concisely observed (Figure 2B-2D).

For *CD4* gene expression, the overall *CD4* gene expression in fish larvae at 21-42 DAYC immunized with FKV-SA significantly differed compared with control larvae ($P < 0.05$) (Figure 2E-

2H). The overall vaccination periods showed significant differences in 21, 28, and 35 DAYC larvae at 336 hav ($P < 0.05$) (Figure 2E-2G), while 42 DAYC larvae showed significantly higher CD4 gene expression at 168 and 336 hav ($P < 0.05$) (Figure 2H).

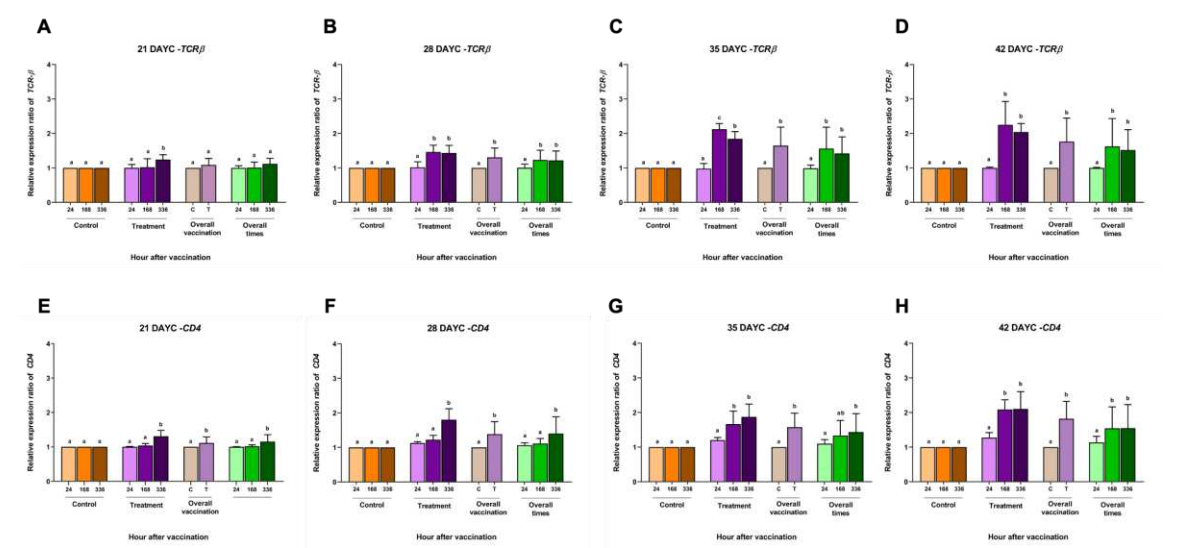


Figure 2. *TCRβ* (A, B, C, D) and *CD4* (E, F, G, H) gene expression at 21, 28, 35 and 42 DAYC. Expression levels were measured by qRT-PCR. Different letters on each bar denote significant differences ($P < 0.05$).

The relative *MHCIIα* gene expression in larvae immunized with FKV-SA 21-42 DAYC showed similarly significant differences compared with the control group at 336 h after vaccination ($P < 0.05$) (Figure 3A-3D). Similarly, significant differences in overall vaccination were observed early at 21 DAYC larvae (Figure 3A). Overall vaccination periods were similarly observed in all groups of 28-42 DAYC larvae after 336 hav of vaccination showed a significant increase only in the group that received the vaccine at 336 hav ($P < 0.05$) (Figure 3A-3D).

The relative *IgM* gene expression was solely found to significantly differ in the 21 DAYC larvae immunized with FKV-SA ($P < 0.05$) (Figure 3E), while 28, 35, and 42 DAYC larvae immunized with FKV-SA were upregulated at 168 and 336 hav ($P < 0.05$) (Figure 3F-3H). Interestingly, all DAYC larvae immunized with FKV-SA demonstrated overall vaccination from 21-42 DAYC larvae ($P < 0.05$) (Figure 3E-3H). Overall, the 21 DAYC larvae significantly differed at 336 hav, while 28, 35, and 42 DAYC were upregulated at 168 and 336 hav ($P < 0.05$) (Figure 3F-3H).

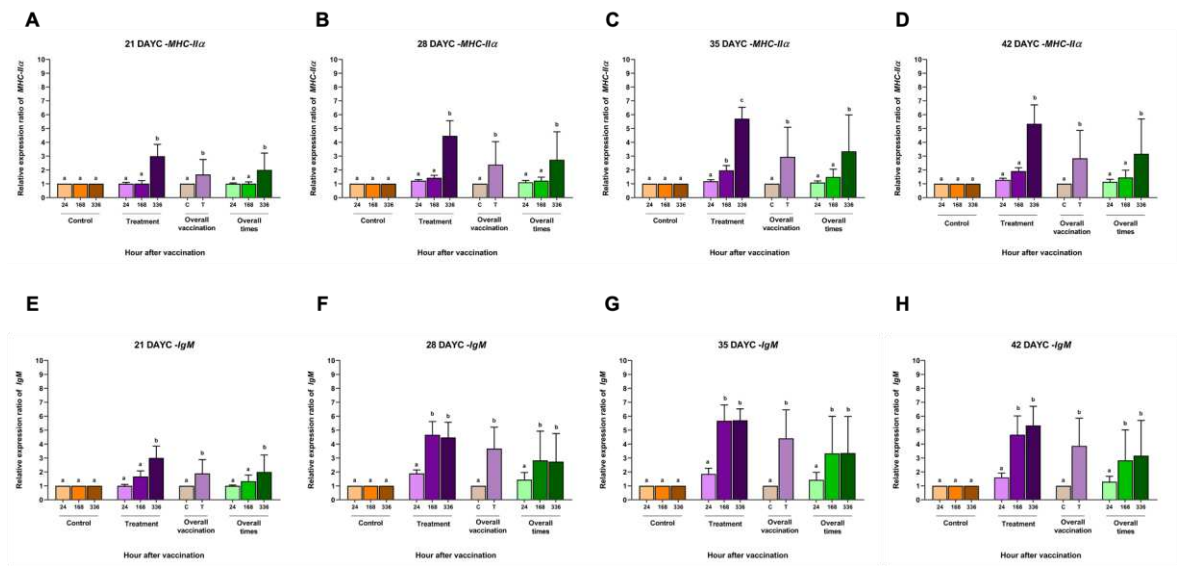


Figure 3. Gene expression analysis of *MHCIIα* (A, B, C, D) and *IgM* (E, F, G, H) at 21, 28, 35 and 42 DAYC. Expression levels were measured by qRT-PCR. Different letters on each bar denote significant differences ($P < 0.05$).

For *IgT* gene expression, larvae at 21 DAYC showed only a significant increase in the larvae group immunized with FKV-SA at 336 hav ($P < 0.05$) (Figure 4A). The vaccinated fish larvae at 28 and 35 DAYC showed significant differences compared with the other groups at 168 and 336 hav ($P < 0.05$) (Figure 4B-4C). However, at 42 DAYC larvae, all immunized fish larvae at all vaccination periods were significantly different from the control larvae ($P < 0.05$) (Figure 4D). Additionally, overall vaccination of the control and vaccinated groups was observed in all DAYC larvae ($P < 0.05$) (Figure 4A-4C). Overall times were significantly observed in 21, 28, and 42 DAYC larvae at 336, 168 and 168 and 336 hav ($P < 0.05$) (Figure 4A-4C). Finally, in the relative *IgD* gene expression, no significant differences between all treatments, overall vaccination, and overall times were observed in 21 and 28 DAYC larvae ($P > 0.05$) (Figure 4E-4F). Significant differences in *IgD* gene expression were observed in the 35 and 42 DAYC larvae at 336, 168, and 336 hav ($P < 0.05$) (Figure 4G and 4H). Significant differences in overall vaccination between the control and vaccinated groups were observed in both 35 and 42 DAYC larvae ($P < 0.05$) (Figure 4G and 4H), while overall vaccination periods were significant in both 35 and 42 DAYC larvae at 336, 168 and 336 hav ($P < 0.05$) (Figure 4G and 4H).

No interaction between periods and vaccination treatments on the expression levels of the genes examined was observed throughout the vaccination periods ($P > 0.05$).

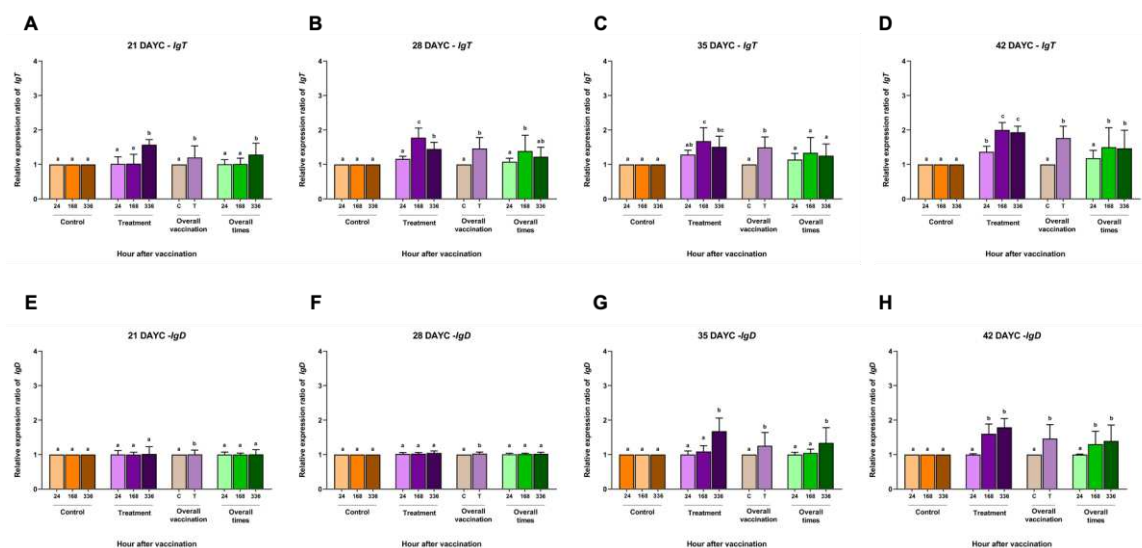


Figure 4. *IgT* (A, B, C, D) and *IgD* (E, F, G, H) gene expression at 21, 28, 35 and 42 DAYC. Expression levels were measured by qRT-PCR. Different letters on each bar denote significant differences ($P < 0.05$).

3.3. The *IgM* distribution in Nile tilapia larvae after immersion vaccination

The distribution of *IgM* in the gills, head kidneys, and intestines of Nile tilapia larvae at 21, 28, and 35 DAYC and 336 hav is shown in Figure 5. At 21 DAYC, the distribution of *IgM* appeared in the tissues of head kidneys and intestines of vaccinated fish (Figure 5G and 5M) compared with the nonvaccinated fish (Figure 5J and 5P). In the gills, a very light signal was observed in the gill lamellae of both groups (Figure 5A and 5D). At 28 and 35 DAYC, fish larvae exposed to FKV-SA strongly demonstrated *IgM* signals in the head kidney (Figure 5H and 5I) and intestines (Figure 5N and 5O), which were more obvious than those in control fish (Figure 5K and 5L, and 5Q and 5R, respectively). In the case of the gills, at 28 and 35 DAYC, higher expression of *IgM* protein was noted with stronger

levels in fish larvae immersed in FKV-SA (Figure 5B and 5C) than in the control groups (Figure 5E and 5F).

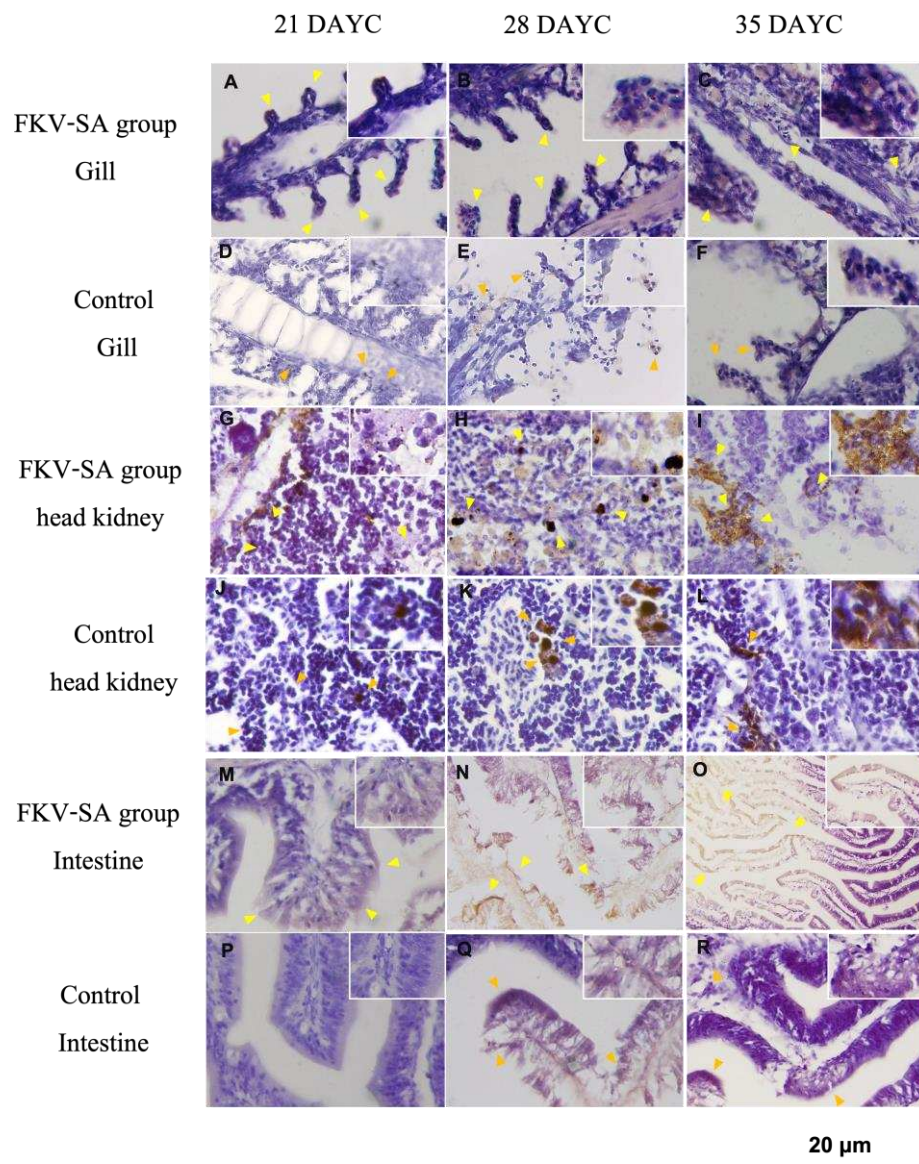


Figure 5. Distribution of IgM in Nile tilapia larvae after immersion FKV-SA vaccination at 336 h based on immunohistochemistry of immunized fish gill FKV-SA group (A, B, C) and control group (D, E, F), head kidney FKV-SA group (G, H, I) and control group (J, K, L), and intestine FKV-SA group (M, N, O) and control group (P, Q, R) at 21, 28 and 35 DAYC.

3.4. Weight and length measurements

The effects of FKV-SA on the weight and total length of Nile tilapia larvae were investigated during vaccination trials. At 336 h, the weights of fish larvae in the control groups at 1, 7, 14, 21, 28, 35, and 42 DAYC were 0.036 ± 0.022 , 0.052 ± 0.011 , 0.098 ± 0.027 , 0.320 ± 0.122 , 0.540 ± 0.220 , 0.620 ± 0.134 , and 0.750 ± 0.123 g, respectively. In contrast, in the vaccinated groups, the weights of fish larvae in the control groups at 1, 7, 14, 21, 28, 35, and 42 DAYC were 0.038 ± 0.027 , 0.059 ± 0.021 , 0.103 ± 0.014 , 0.450 ± 0.231 , 0.641 ± 0.194 , 0.730 ± 0.207 , and 0.821 ± 0.190 g, respectively. The lengths of fish larvae in the control groups at 1, 7, 14, 21, 28, 35, and 42 DAYC were 1.10 ± 0.6 , 1.29 ± 0.2 , 1.39 ± 0.1 , 1.49 ± 0.2 , 1.73 ± 0.1 , 1.83 ± 0.2 , and 1.94 ± 0.2 cm, respectively. Furthermore, the values for the fish larvae in the vaccinated groups at 1, 7, 14, 21, 28, 35, and 42 DAYC were 1.30 ± 0.1 , 1.42 ± 0.1 , $1.59 \pm$

0.1, 1.67 ± 0.2 , 1.84 ± 0.1 , 1.97 ± 0.2 , and 2.18 ± 0.2 cm, respectively. However, the growth indices had no statistically significant changes (Supplementary Table S1 and Supplementary Table S2).

3.5. Lethal concentration (LC_{50}) of *S. agalactiae*

On day 8, the cumulative percent mortality of Nile tilapia larvae at concentrations 1×10^4 , 1×10^5 , 1×10^6 , 1×10^7 , and 1×10^8 CFU/mL was 31.11 ± 3.33 , 35.56 ± 6.94 , 47.78 ± 5.09 , 74.44 ± 8.39 , and $82.22 \pm 3.33\%$, respectively (Figure 6). Based on the Probit analysis, the LC_{50} of *S. agalactiae* in Nile tilapia larvae was $3.44 \pm 1.26 \times 10^5$ CFU/mL. This concentration was further used for the challenge test.

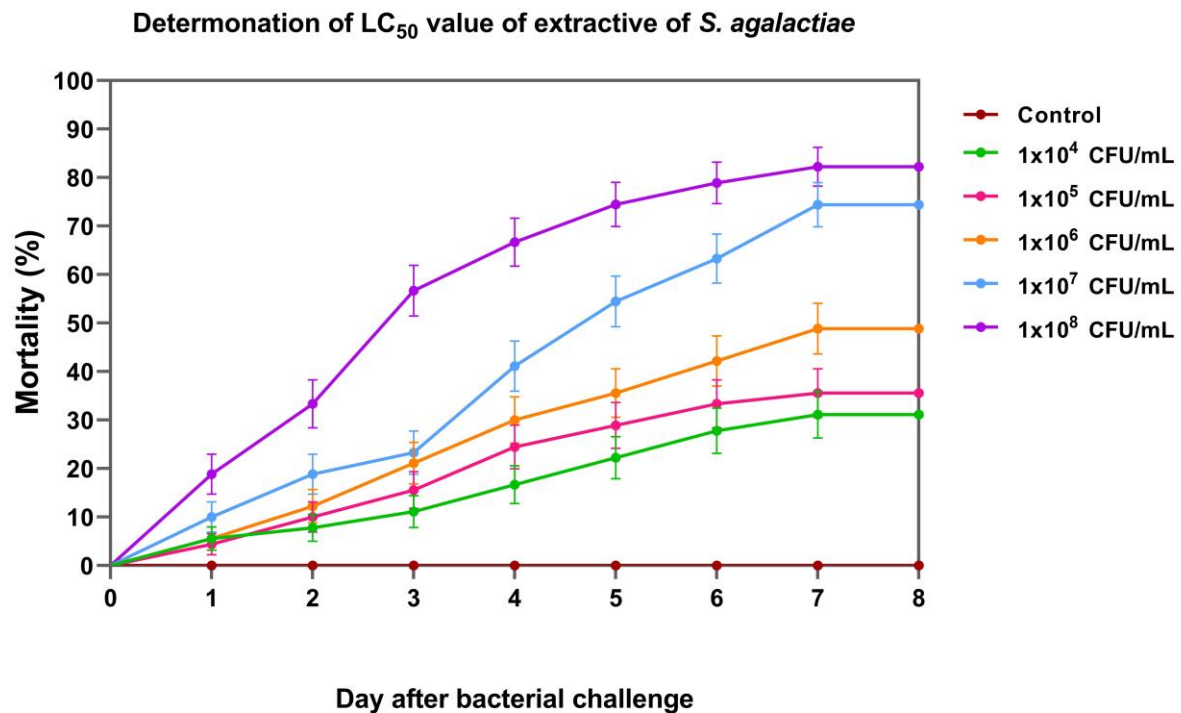


Figure 6. Cumulative mortality of Nile tilapia larvae exposed to different concentrations of *S. agalactiae* for median lethal concentration (LC_{50}) analysis.

3.6. Survival analysis and RPS of Nile tilapia challenged with *S. agalactiae*

In this experiment, 21, 28, 35, and 42 DAYC Nile tilapia larvae immunized with FKV-SA for 24, 168, and 336 hav were challenged with viable *S. agalactiae* to test the efficacy of vaccines with various exposure times. The survival rates of vaccinated Nile tilapia larvae after challenge with *S. agalactiae* are shown in Figure 7A-7H. In fish 21 DAYC larvae, fish immunized with only FKV-SA for 336 hav significantly differed from control fish larvae ($P < 0.05$) by 58.67 ± 2.31 and $42.00 \pm 3.61\%$, respectively (Figure 7A). In 28 DAYC larvae, significant differences in survival between vaccinated and unvaccinated groups were observed at 168 hav (66.00 ± 4.58 and $40.61 \pm 2.08\%$) and 336 hav (62.00 ± 2.00 and $44.00 \pm 2.65\%$) after vaccination ($P < 0.05$) (Figure 7B). Similar results were observed in the 35 and 42 DAYC larvae. In 35 DAYC larvae, vaccinated fish showed significant differences at 168 and 336 hav with survival rates of 65.33 ± 2.31 and $62.00 \pm 5.57\%$, respectively, compared with the control by 40.67 ± 1.53 and $44.67 \pm 2.08\%$, respectively ($P < 0.05$) (Figure 7C). Similarly, in 42 DAYC larvae, vaccinated fish showed significant differences at 168 and 336 hav with survival rates of 68.67 ± 3.79 and $62.67 \pm 2.52\%$, respectively, compared with the control by 42.00 ± 4.00 and $46.00 \pm 2.65\%$, respectively ($P < 0.05$) (Figure 7D). Moreover, the RPSs of fish larvae immunized with FKV-SA at various exposure times were also calculated. Larvae at 21 and 28 DAYC, and at 24, 168, and 336 hav did not significantly differ between exposure times ($P > 0.05$) (Figure 7E and 7F). However, in 35 and 42 DAYC larvae, it was found that fish larvae immunized with FKV-SA for 168 hav showed higher

significant differences in RPSs than those immunized at 24 and 336 h, with the highest RPSs of 41.53 ± 7.61 and $45.54 \pm 13.44\%$, respectively ($P < 0.05$) (Figure 7G and 7H).

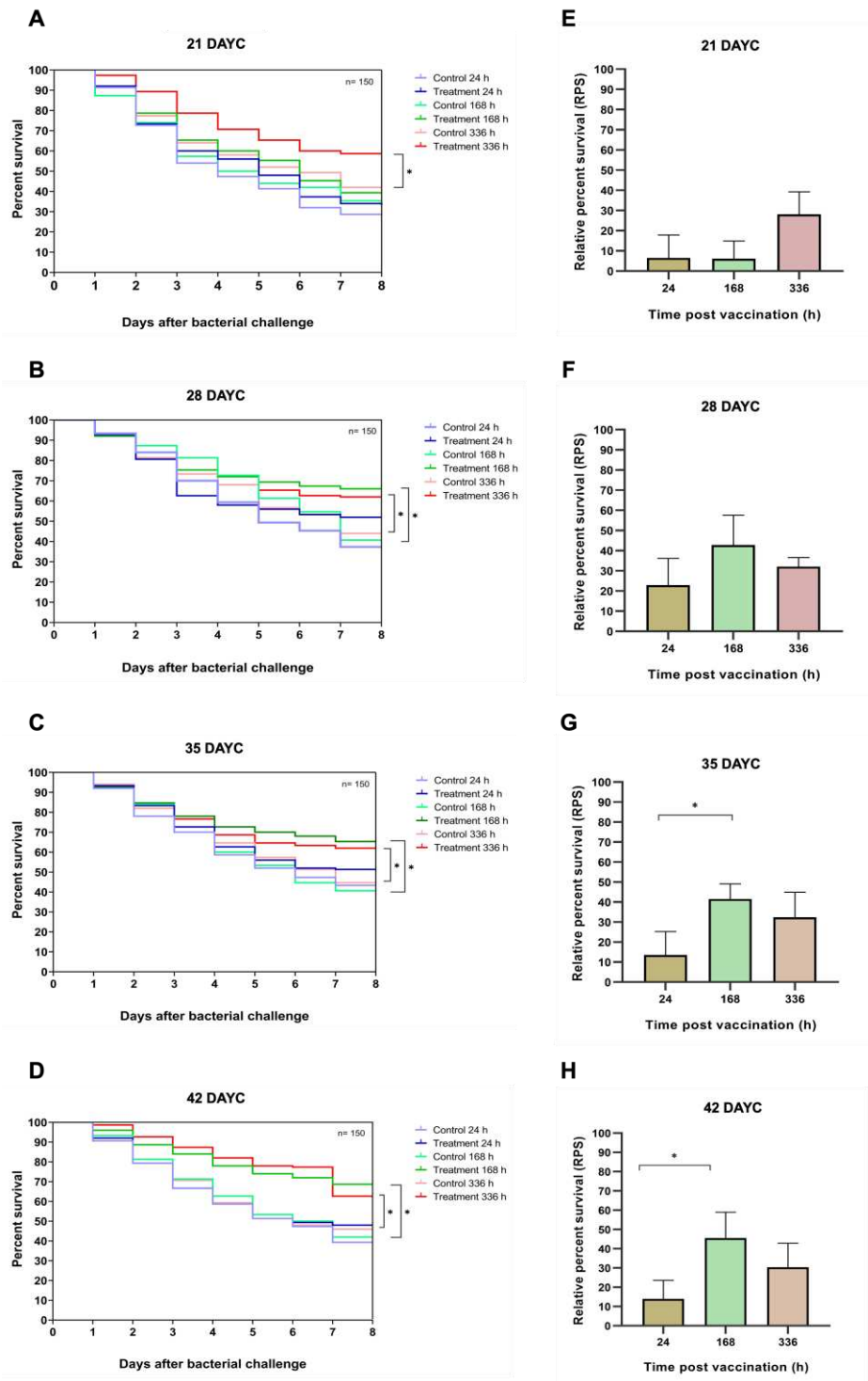


Figure 7. Data and survival plots were performed using the Kaplan-Meier method (A-D). Survival analysis and relative percent survival (RPS) of Nile tilapia FKV-SA vaccinated after challenges with *S. agalactiae* at 21, 28, 35 and 42 DAYC (E-H). The levels of statistical significance between the control and treatment groups are indicated by * ($P < 0.05$).

4. Discussion

Streptococcosis caused by *S. agalactiae* is classified as a major constraint that hampers Nile tilapia aquaculture worldwide [24-27]. Vaccines are one of the first measures selected to overcome this problem. Based on the current information, various research studies have been reported to develop effective vaccines against *S. agalactiae* infection in Nile tilapia [13, 28-31]. Even though the obtained results of most developed vaccines were very effective at the laboratory [32-34] and field trial scales [35-37], most of those reports have relied on injection methods [38-40]. This vaccination route was subsequently found to be impractical and unacceptable for field application. Therefore, a more practical vaccination is needed. Oral and immersion vaccinations are the most important vaccines of interest. Immersion vaccination has several advantages over injection and oral administration, such as cost-effectiveness, low-stress responses, labor, and small size of fish needs, and numerous fish in each vaccination can be conducted [17,41,42]. However, some crucial disadvantages remain, especially the use of high amounts of vaccines, feeble immune responses, and less protection in small fish [43,44]. Additionally, the most fragile aspect of this strategy is that the most effective age and size for immersion vaccination of fish are obscured in many fish species, including Nile tilapia.

Similar to other higher vertebrates, early immune responses in fish can be studied by examining the development of immune-related organs. Unfortunately, very few studies have shown early immune responses in Nile tilapia.

Previously, Cao et al. (2017) [45] investigated the development and differentiation mechanisms of the thymus gland using specific hybridization of the *RAG-1* gene in the early larval stage to the reproductive stage at 2 days postfertilization (dpf) through 1 year of age. Importantly, it has been observed that the thymus gland of tilapia starts developing at an early stage, approximately 2 and 5 dpf; the thymus gland increases in size according to the number of thymic cells, while at 7 and 10 dpf, the thymus gland shows a noticeable increase in size and a significant increase in the number of thymic cells. In addition, the thymus formed as a bulge on the surface dorsolateral to the pharyngeal cavity at those stages [46]. Furthermore, variations in the Nile tilapia embryo and larva development rates were found, and a prominent thymus gland was present at 100 h. Hematopoietic tissue develops near the pronephros during early larval development [47]. In the case of Nile tilapia broodstock and larval immunization, Pasaribu et al. (2018) evaluated the efficacy of maternal transfer and offspring protection resulting from immunization with monovalent and bivalent vaccines (Biv). The relative percent survival of the Biv group revealed significantly enhanced disease resistance against *S. agalactiae*, *A. hydrophila*, and their coinfection [48]. However, these results are too few to effectively apply to Nile tilapia vaccination. Therefore, our present study focused on assessing specific IgM antibody responses, analyzing specific-immune gene expression under vaccination conditions, and evaluating vaccine efficacy through challenge tests.

In the current study, we found the presence of IgM molecules by ELISA and immunohistochemistry and the expression patterns of immune-related genes, including *IgM*, *IgT*, *IgD*, *MHCII α* , *TCR β* , and *CD4*, which are great indicators of specific immune systems [49]. The results showed that larvae at 21 DAYC (0.108 ± 0.110 g) constitute the earliest stage to effectively express specific immune components post-immersion immunization with inactivated *S. agalactiae* vaccine, which suggests that it is an immunocompetent stage of Nile tilapia.

Fish in tropical areas generally reach the immunocompetent stage faster than fish in subtropical and temperate zones [50]. It takes approximately 28-30 days after hatching to develop full immunocompetence. This is often marked by the appearance of the thymus, a primary lymphoid organ, and other immune-related cells and organs [51]. In Nile tilapia, the thymus is known to initially develop as early as two days after hatching but remains very small. As fish move into the immunocompetent stage, the thymus continues to grow progressively during the juvenile phase and then decreases in size as the animals age and their spines develop [45]. The results of this research and the findings from previous studies consistently suggest that the age range of 28-30 days after hatching is a critical time when the immune system reaches a stage of full competence [51]. This stage signifies the transition from an immature to a fully functional immune system. However, the above information is insufficient to clearly describe the immunocompetent stage in Nile tilapia.

In this study, the fish at 28-42 DAYC (0.33-0.58 g) showed a remarkably rapid response compared to those at 21 DAYC (0.11 g). It is possible that the fish in these age groups had more developed immune systems that enabled them to respond more effectively to the vaccines than those at 21 DAYC. Compared to other studies in which fish were exposed to antigens via immersion vaccination, Nile tilapia of various ages or weights (sizes) may produce different specific antibodies. For example, in rainbow trout (*Oncorhynchus mykiss*), an increase in the levels of antibodies to *Vibrio anguillarum* was observed in juvenile fish weighing approximately 0.14 g or 2 weeks after hatching [52]. In the case of young Nile tilapia weighing 0.1 ± 0.01 g, they could efficiently form antibodies against *S. agalactiae* [53]. A significant increase in antibody levels was observed in channel catfish (*Ictalurus punctatus*) after vaccination with *Edwardsiella ictaluri* compared with the control group. The first significant increase was observed in fish aged 4 weeks with an average weight of 0.085 g [54]. In Asian seabass (*L. calcarifer*) at 35 and 42 dph with weights of approximately 0.10 ± 0.03 and 0.25 ± 0.14 g, respectively, a statistically significant increase in specific IgM levels against *S. iniae* was observed after administration of *S. iniae* heat-killed vaccine after 7 days [55].

The expression of genes associated with the specific immune response of Nile tilapia larvae at 21, 28, 35, and 42 DAYC and at 24, 168, and 336 h after stimulation with FKV-SA was investigated. Normally, TCR β is the receptor protein of T lymphocytes, while CD4 is essential in verifying the accuracy of binding between TCR $\alpha\beta$ -epitope- MHCII $\alpha\beta$. Moreover, MHCII $\alpha\beta$ presents the structure of epitopes, which are partial components of pathogens or antigens, and their bond to T cells or B cells to stimulate these cells to produce cytokines. This stimulation involves T cells, B cells, or APCs, which trigger proliferation and differentiation processes to become memory cells, such as memory T or B cells, to remember pathogens or antigens of the same type or plasma B cells for the production of immunoglobulins (Ig) to remember specific diseases and antigens [56-58]. Based on this study, it was found that the expression of TCR β , CD4, and MHCII α genes in Nile tilapia fry at 21, 28, 35, and 42 DAYC that had received FKV-SA showed complete expression at all time intervals, especially after 336 hav. This expression was significantly higher than that of the unvaccinated control group. The immunoglobulin (Ig) types found in Nile tilapia with bony structures include IgM, IgT, and IgD [59]. The secreted immunoglobulin molecule (sIg) is critical in processes such as neutralization, opsonization, antibody-dependent cytotoxicity, and the complement activation system. These components maximize the efficiency of disease and antigen control or the immune response [60]. We found that the expression of IgM and IgT genes in Nile tilapia larvae at 21, 28, 35, and 42 DAYC immunized with FKV-SA showed higher expression at all time intervals compared with unvaccinated fish, especially after 336 hav. In contrast, the IgD gene was specifically expressed in tilapia at 35 and 42 DAYC, with higher expression observed in the vaccinated group at 336 hav. The results suggest that IgM and IgT of Nile tilapia are crucial Igs in the response and protection of fish to invading pathogens at early stages. Although IgD was found to be a late response, the results suggest that it is a non-readiness molecule during the early stage of fish development. However, anti-IgT and anti-IgD development is needed to further understand this obscured phenomenon.

Additionally, the vaccine was effective in upregulating several immune-related genes, such as TCR β , CD4, MHCII α , IgM, IgT and IgD, in immune-related tissues, including the head kidney, spleen, PBLs, and gills, which is well supported by previous studies and other publications [61, 62]. For example, in Asian seabass (*Lates calcarifer*) at 35 and 42-day post-hatching (dph) at 7 days after vaccination, there was a statistically significant increase in the expression of CD4, MHCII α , IgM, IgT, and IgD genes [55]. In juvenile Nile tilapia, various nanovaccines were used to treat francisellosis and columnaris diseases by immersion. It was observed that the expressions of TCR β , CD4, MHCII α , IgM, and IgT genes significantly increased in the head kidney, spleen, gills, and peripheral blood leukocytes (PBLs) 8 weeks after the first vaccine administration [19].

After immersion vaccination in tilapia of different ages, based on immunohistochemistry at 21, 28, and 35 DAYC, it was shown that IgM was distributed in different organs. The most densely distributed IgM was found in the head kidney and intestine. The organ with the highest IgM reaction was the head kidney, which is consistent with the results observed in *Takifugu rubripes* in all organs, with higher expressions in the primary lymphoid organs such as the head kidney and spleen [63]. In

Pseudosciaena crocea and *Larimichthys crocea*, significant IgM heavy chain gene expression was detected in the spleen, peripheral blood, and head kidney [64, 65]. Furthermore, using the immunohistochemistry technique in Nile tilapia after receiving a killed *S. agalactiae* vaccine, it was found that the distribution of IgM was observed across various organs of the fish, with a prominent presence in the head kidney [53].

The above-obtained information strongly supports the phenomena found in the challenge test experiment. When the 21-42 DAYC larvae were immunized with FKV-SA, 21 DAYC showed significantly higher differences in survival than the control at only 336 h after vaccination, which was different from the information found in 28-42 DAYC larvae. In these groups, fish larvae immunized with FKV-SA showed significantly higher differences than unvaccinated fish against viable *S. agalactiae* at 168 and 336 h, which was 7 days earlier than what was observed in 21 DAYC larvae. This indicated that the optimal period for initial vaccination in Nile tilapia larvae is 28-35 DAYC larvae.

This study revealed that administering FKV-SA to Nile tilapia larvae from 21 DAYC onward can effectively stimulate a specific and targeted immune response against *S. agalactiae* infection. This success can be attributed to the role of both primary and secondary lymphoid organs and tissues, particularly the various types of mucosa-associated lymphoid tissues (MALTs). These tissues are characterized by their mucous structure and their involvement in the general immune response. Fish possess mucous tissues that cover every surface of their bodies, and thus, the structure of these mucous tissues surrounding the fish body is crucial in forming a specific immune system capable of detecting environmental disturbances. They serve as the first line of defense against disease pathogens, enabling fish to interact with the external environment that surrounds them. In particular, gill-associated lymphoid tissue (GALT) is essential. Additionally, lymphoid tissue in the skin, referred to as skin-associated lymphoid tissue (SALT), has also been reported with components related to immunity. These findings have confirmed that these tissues can indeed trigger responses against diseases and antigens [66-69].

Although the vaccination method used in this study involved immersion, there are reports indicating that immersion vaccination can stimulate the immune system's activity in GALT within the gut. It has been reported that B cells, particularly IgM and IgT B cells, are found in the lamina propria (LP) layer of the intestine, which is a part of the gut mucosa [70,71]. Additionally, due to the behavior of freshwater fish not drinking water to regulate their body's salt and mineral balance (osmoregulation) but rather engaging in sipping behaviors, there is a chance that the vaccine, dissolved in water, can enter the digestive system. This can lead to stimulation of the GALT in the gastrointestinal tract of the fish, similar to how it happens with immersion vaccines. The fish digestive system is considered a crucial area that is directly interconnected with the external environment through ingestion. This process provides a pathway for potential disease pathogens and antigens to enter. Moreover, it serves as a site for triggering specific immune responses, especially in the mucosal immune system, such as GALT within the intestine. GALT is crucial in maintaining gut equilibrium and acts as a defense mechanism against various diseases and foreign agents [72].

Based on the study of disease resistance using *S. agalactiae* in Nile tilapia larvae at 21, 28, 35, and 42 DAYC and at 24, 168, and 336 h postvaccination with the FKV-SA vaccine, it was observed that the postvaccination survival rates between the vaccinated and control groups showed significant differences starting at 21 DAYC after 336 h of vaccination. However, for 28, 35, and 42 DAYC, the vaccinated fish exhibited significant disease resistance compared to the control group after 168 and 336 h of vaccination across all age groups. Furthermore, the RPS values were found to differ significantly between vaccinated and nonvaccinated fish in each age group (21-42 days). However, the RPS values obtained ranged from relatively low values of 28.14 to 45.54. This suggested that while the fish in these age ranges have developed specific immune responses to protect against and resist *S. agalactiae* infection, the effectiveness of the immune response is not very high. Generally, a level of 60 or higher is considered acceptable [73].

Based on this, more suitable vaccine formulations should be developed. Currently, there are reports suggesting that the effectiveness of immune responses can be improved through vaccination

methods, such as immersion, which enhances the overall immune response. For instance, the utilization of mucoadhesive cationic lipid-based nanoencapsulation for vaccines to prevent columnaris disease in Asian seabass (*Lates calcarifer*) through immersion resulted in a survival rate of $72.50 \pm 3.54\%$ and the highest RPS recorded at 62.07 ± 4.87 . Similarly, the application of nanovaccines to combat francisellosis and columnaris diseases through immersion in juvenile Nile tilapia has exhibited survival rates ranging from 65.83 to 72.50% and RPS values ranging from 52.87 to 62.07 [3,19]. Furthermore, the utilization of mucoadhesive polymers to enhance the effectiveness of inactivated vaccines in preventing columnaris diseases caused by *F. columnare* infection in red tilapia has shown a maximum survival rate and RPS of 83% and 81, respectively [74]. These results showcase an additional approach to enhance the efficacy of inactivated vaccines for preventing harmful bacterial diseases through immersion.

Finally, the results from the current study demonstrated that specific immune responses start developing in Nile tilapia larvae from 21 DAYC onwards. The knowledge gained from this study is crucial and can be applied to the development of vaccines in small Nile tilapia, enabling them to effectively prevent streptococcosis, a disease commonly occurring in the fish farming industry.

5. Conclusions

Administration of the *S. agalactiae* vaccine (FKV-SA) to Nile tilapia larvae by immersion can initially induce specific IgM and expression of targeted immune-related genes in the fish as early as 21 days after yolk sac collapse (DAYC). It is possible that the complete developmental stage of the fish reaches maturity at approximately 21 DAYC, indicating the critical period when Nile tilapia larvae develop immunocompetence. Based on key specific immune gene expression data and challenge tests against viable *S. agalactiae*, we recommend the optimal vaccination period for Nile tilapia larvae to be between 28 and 35 DAYC. However, the RPS of vaccinated larvae in various DAYCs is relatively low. This result suggests that although vaccination can elicit a crucial immune response in fish, protection against disease is relatively inefficient. This finding could have been obtained due to several factors, such as low immunogenicity of the FKV-SA form, low antigen uptake of immersion administration, or immature cell-mediated immunity (not investigated in this study), in the current study, which requires further research and development in the future. Nonetheless, the results of this study provide important baseline knowledge that can be used to develop effective vaccination methods to reduce constraints and losses in Nile tilapia production caused by severe and highly damaging disease outbreaks. Early vaccination may contribute to sustainable, non-antibiotic farming of Nile tilapia in the near future.

Supplementary Materials: The following supporting information can be downloaded at www.mdpi.com/xxx/s1, Table S1: Growth performance by weight of Nile tilapia larvae was investigated during vaccination trials; Table S2: Growth performance by total length of Nile tilapia larvae was investigated during vaccination trials.

Author Contributions: B.K.; Visualization, Data Curation, Formal Analysis, Investigation, Writing - Original Draft. A.B.; Conceptualization, Methodology, Formal Analysis, Writing - Review and Editing. S.C.; Conceptualization, Writing - Review and Editing. P.K.; Conceptualization, Methodology. H.T.D.; Conceptualization, Methodology, Writing - Review and Editing. S.S.; Conceptualization, Funding acquisition, Writing - Review, Funding acquisition, and Editing. P.S.; Conceptualization, Resources, Supervision, Funding acquisition, Writing - Review and Editing.

Funding: This research was funded by Kasetsart University through the Graduate School Fellowship Program and BIOTEC Fellow's Research Grant (P-21-51690).

Data Availability Statement: Suggested Data Availability Statements are available in the section "MDPI Research Data Policies" at <https://www.mdpi.com/ethics>.

Acknowledgments: The authors would like to thank Kornsunee Phiwsaiya for the technical assistance provided.

Conflicts of Interest: The authors declare no conflict of interest.

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