

Article

Effects of Dietary *Enterococcus faecalis* YFI-G720 on the Growth, Immunity, Serum Biochemical, Intestinal Morphology, Intestinal Microbiota, and Disease Resistance of Crucian Carp (*Carassius auratus*)

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Abstract: Diseases of crucian carp (*Carassius auratus*) are closely related to intestinal parameters. *Enterococcus faecalis* has strong colonization ability in the intestinal tract, and produces natural antibiotics, bacteriocin, and other bacteriostatic substances, which can effectively inhibit some pathogenic bacteria and improve the intestinal microenvironment. This study aimed to assess the effects of *E. faecalis* YFI-G720 which was isolated from the intestinal of crucian carp on the growth, immunity, intestinal health, and disease resistance of crucian carp. Fish (48.16 ± 0.55 g) were fed four diets, commercial diet or diet containing *E. faecalis* at 10^5 CFU/g (EF1), 10^6 CFU/g (EF2), or 10^7 CFU/g (EF3) for 28 days. The results showed that supplementation of *E. faecalis* significantly improved the weight gain ratio (WGR) and the specific growth rate (SGR) compared with control group ($p < 0.05$). Intestinal mucosal epithelial cells in EF2 were intact and normal, but there was obvious vacuolation in CG. Compared with CG, serum C3 and IgM in EF2 were significantly increased at the end of the experiment ($p < 0.05$), and serum alkaline phosphatase was significantly higher in all experimental groups ($p < 0.05$). Among studied immune-related genes, expression was detected by qPCR, C3, IgM, and IL-1 β were upregulated in all experimental groups to varying degrees from 14 days, with highest expression in EF2 at 28 days. Intestinal microbiota structure analyzed through high-throughput sequencing, and the results showed that the relative abundance of *Aeromonas* and *Acinetobacter* decreased while *Cetobacterium* increased in all experimental groups, with the greatest changes in EF2. Challenge tests showed that fish fed *E. faecalis* were more resistant to *Aeromonas veronii* ($p < 0.05$). In conclusion, dietary *E. faecalis* YFI-G720 at 10^6 CFU/g can improve the health status, immune parameters, intestinal microbiota composition, and disease resistance of crucian carp.



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Keywords: crucian carp; *Enterococcus faecalis*; growth; immunity; serum parameters; intestinal microbiota; intestinal morphology

1. Introduction

Crucian carp (*Carassius auratus*), an omnivorous and bottom-feeding freshwater fish, is one of the main aquaculture species in China [1,2]. In 2020, total production of crucian carp in China was 2.75 million tons, which constituted about 8.9% of the total freshwater fish aquaculture production [3]. Crucian carp meat contains 13% protein, 11% fat, and plenty of minerals, hence it is appreciated by consumers [4]. However, with the increasing development and intensification of aquaculture, diseases caused by pathogens affecting this species are gradually increasing, such as *Aeromonas veronii*, *Aeromonas hydrophila*, and *Cyprinid herpesvirus 2* (CyHV-2) [5–7]. These pathogens mainly spread to other tissues and organs after infecting the intestinal tract. Therefore, the occurrence of diseases is closely related to intestinal health, especially the microbial composition of the intestinal [8,9]. The

(species) diversity and stability of the intestinal microbial are important factors affecting host health [10,11]. Therefore, for the crucian carp industry, it is important to maintaining a balance of intestinal microbiota via good health management.

Probiotics have been proven to promote intestinal health, improve feed digestibility and absorption rate, and enhance disease resistance by regulating the intestinal microbiota [12,13]. *Enterococcus faecalis*, a facultative anaerobic Gram-positive bacterium belonging to the *Enterococcaceae* within *Lactobacillus* and *Enterococcus*, is one of the major intestinal microbiota components in humans and animals [14–16]. *E. faecalis* has strong tolerance to the environment and colonization ability in the intestinal tract that can help to form a barrier of lactic acid bacteria, preventing the invasion of foreign pathogens and viruses [17]. Additionally, its secondary metabolites—including organic acids, hydrogen peroxide, extracellular polysaccharides, and other substances—can inhibit the growth and reproduction of the various pathogenic bacteria [16,18]. *E. faecalis* can also facilitate the digestion and absorption of feed, enhance the activity of macrophages, and promote the immune responses of the host [19,20]. Studies have shown that dietary supplementation with 10^7 – 10^9 CFU/g *E. faecalis* can include the growth, immunity, and disease resistance of rainbow trout (*Oncorhynchus mykiss* Walbaum), Javanese carp (*Puntius gonionotus* Bleeker 1850), and mud crab (*Scylla paramamosain*) [19–21]. Thus, *E. faecalis* provides an important relevance for the development of probiotics in aquatic products.

A number of studies have verified the beneficial effects of probiotic bacteria such as *Bacillus* and *Lactobacillus* on crucian carp [22,23]. However, data regarding the effects of *E. faecalis* on the growth, immunity, and intestinal health of crucian carp are limited. *E. faecalis* YFI-G720 was isolated from the intestinal tract of crucian carp, and preliminary experiments showed that it could improve the intestinal microbiota of crucian carp. Therefore, the purpose of this study was to examine the effects of *E. faecalis* YFI-G720 supplementation of diet on growth performance, immune capacity, intestinal morphology, and the intestinal microbiota of crucian carp, as well as providing a reference for the application of probiotics in the crucian carp industry.

2. Materials and Methods

2.1. Experimental Diets

Experimental diets comprised puffed pellet feed which supplied by Tongwei Co., Ltd. (Chengdu, China). *E. faecalis* YFI-G720 stored at the China Center for type Culture Collection (CCTCC), Wuhan University, under preservation number CCTCC M2021312. *E. faecalis* was cultured overnight on brain heart infusion plates (BHI, Qingdao biological technology co., Ltd., Qingdao, China) at 28 °C and resuspended in sterilized phosphate-buffered saline (PBS). Colony-forming units (CFU) per mL of *E. faecalis* culture was determined by the plate dilution counting method. The *E. faecalis* cells were resuspended and sprayed on the commercial basal feed at 10^5 CFU/g, 10^6 CFU/g, and 10^7 CFU/g of diet [18], followed by thorough mixing. Feeds were vacuum dried at 30 °C overnight, stored in individual airtight containers at 4 °C, and produced every 3 days to maintain the cell count of *E. faecalis*.

2.2. Laboratory Fish and Rearing

The experiment was carried out in the Fish Disease Laboratory of the Yangtze River Fisheries Research Institute. Experimental crucian carp were obtained from a local farm in Wuhan, Hubei, China, and we randomly sampled for parasite observation, bacterial isolation, and CyHV-2 detection, and the results showed negative. Then crucian carp were acclimated in 300 L freshwater aquaculture tanks for 15 days. Next, 600 fish with a mean initial body weight of 48.16 ± 0.55 g were randomly assigned into four groups (three tanks per group, 50 fish per tank). The control group was fed with commercial basal diet, while EF1, EF2, and EF3 groups were fed *E. faecalis* diet supplemented at 10^5 CFU/g, 10^6 CFU/g, and 10^7 CFU/g, respectively. Fish were fed twice daily at 08:30 morning and 17:00 afternoon a day with the feeding rate of 3–5% of body weight. A quarter of the tank water was replaced every day. During the trial (4 weeks), the tank water

temperature was maintained at 23–25 °C, pH 7.5–8.0, dissolved oxygen > 7.0 mg/L and flow rate > 4 mL/s. All experimental procedures were conducted according to the guidelines of the appropriate Animal Experimental Ethical Inspection of Laboratory Animal Center, Yangtze River Fisheries Research Institute, Chinese Academy of Fishery Sciences.

2.3. Growth Performance

Body weight of fish was measured at the beginning and the end of the experiment. Growth parameters were calculated according to the following formulae to measure the growth performance of crucian carp: WGR (weight gain ratio, %) = $100 \times (\text{final average body weight} - \text{initial average body weight}) / \text{initial body weight}$; SGR (specific growth rate, %/day) = $100 \times [\ln(\text{final average body weight}) - \ln(\text{initial average body weight})] / \text{days}$; CF (condition factor, %) = $100 \times \text{final body weight} / [\text{body length (cm)}]^3$; Survival rate (%) = $100 \times \text{final number of survived fish} / \text{initial number of fish}$.

2.4. Sample Collection

After 7, 14, 21, and 28 days of starting feed with special diet, three crucian carp from each tank were randomly selected, 500 µL blood was collected from the caudal vein of each fish with a 1 mL syringe and placed in plastic Eppendorf tubes containing anticoagulant solution (heparin). The tubes were kept at 4 °C overnight, centrifuged at $4000 \times g$ for 10 min, and the resulting serum was stored at –80 °C for serum biochemical index analysis. Meanwhile, spleen tissue was placed in a RNase-free centrifuge tubes containing 200 µL TRIzol reagent (Invitrogen, Carlsbad, CA, USA) and stored at –80 °C for immunity gene analysis.

At the end of the experiment, three crucian carp from each tank were randomly collected and aseptically sacrificed in an ice bath. The anterior intestines were fixed in neutral 4% paraformaldehyde for analysis of intestinal morphology. Intestinal tissue was collected, frozen rapidly in liquid nitrogen, and stored at –80 °C for intestinal microbiota analyses.

2.5. Serum Biochemical Analysis

Complement 3 (C3), immunoglobulin M (IgM), triglyceride (TG), total cholesterol (TCHO), alkaline phosphatase (AKP), and aspartate amino transferase (AST) in serum were determined by Olympus600 automatic biochemical analyzer (Olympus, Tokyo, Japan) using a commercial kit (Guangzhou Dibao Medical Products Co., Ltd., Guangzhou, China).

2.6. Real-Time PCR Analysis of Immune-Related Genes

Immunity gene expression levels in spleen tissue of crucian carp were examined by quantitative fluorescence real-time PCR (qPCR) method. Total RNA was isolated using the TRIzol reagent (Invitrogen, Carlsbad, CA, USA) and the residual trace of DNA was removed using Recombinant DNase I (RNase-free, Takara, Dalian, China). Complementary DNA (cDNA) was synthesized using a cDNA Synthesis SuperMix Kit (TransGen Biotech, Beijing, China). All qPCR experiments were performed using a SYBR Green Premix Ex Taq Kit (Takara, Taejin, Japan) by Rotor-Gene 6000 Real-time PCR System (Qiagen, Dusseldorf, Germany). Thermal cycling included denaturation at 95 °C for 5 min, followed by 30 cycles at 95 °C for 20 s, 56 °C for 20 s, 72 °C for 20 s, and 72 °C for 20 s. The primers for qPCR are shown in Table 1. The IgM gene sequence was found on NCBI (MK272741.1), and the primer was designed by primer 5. All experiments were repeated at least three times, and relative expression levels were calculated using the $2^{-\Delta\Delta C_t}$ method with the β -actin gene used as an internal control gene for cDNA normalization.

Table 1. Sequences of primer pairs used in real-time PCR.

Genes	Primer Sequence (5'–3')	GenBank Number
β -action-F β -action-R	GATGATGAAATTGCCGCACTG ACCGACCATGACGCCCTGATGT	AB039726 [24]
IL-1 β -F IL-1 β -R	GCGCTGCTCAACTTCATCTTG GTGACACATTAAGCGGCTTCA	AJ249137 [24]
C3-F C3-R	AGTGAAATGGTGGGAAGCAGAAAG TACGTATACCGAGACATCGAAGG	KF110786 [25]
IgM-F IgM-R	GTGGAACCTTGATGCCCAAT CATCAGCAAGCCAAGACACAA	MK272741.1

2.7. Histopathology Analysis

Anterior intestine tissues were fixed in 4% paraformaldehyde for 24 h, dehydrated in a sequential ethanol series of alcohol (50–95%) and embedded in paraffin. Tissue blocks were sectioned (5 μ m thick) and stained with hematoxylin and eosin (H&E). The intestinal structure was assessed and photographed under an Olympus BX53 microscope (Olympus, Tokyo, Japan).

2.8. Genomic DNA Extraction and 16S rRNA Gene Sequencing

Bacterial genomic DNA was extracted using a Bacterial DNA Kit (Omega Biotek, Winooski, VT, USA) following manufacturer's instructions. PCR amplification of the bacterial V4–V5 region of the 16S rRNA gene was performed in a 50 μ L reaction. The reaction contained 25 μ L of Hot Start Taq 2 \times Master Mix (New England BioLabs Inc., Ipswich, MA, USA), 2 μ L of template DNA, 2 μ L of each primer (338F, ACTCCTACGGGAGGCAGCA; 806R, GGACTACHVGGGTWTCTAAT [26]), and deionized water. Thermal cycling comprised an initial denaturation at 95 $^{\circ}$ C for 2 min, followed by 30 cycles at 95 $^{\circ}$ C for 15 s, 55 $^{\circ}$ C for 30 s, and 72 $^{\circ}$ C for 30 s, and a final extension at 72 $^{\circ}$ C for 5 min. After checking the quality by 1% agarose gel electrophoresis, samples were sequenced on an Illumina MiSeq PE300 high-throughput sequencing platform. The final effective data were obtained using the UCHIME v4.2 software [27]. Reads were clustered into operational taxonomical units (OTUs) at 97% identity [27]. The abundances of the corresponding OTUs in each group were calculated at the phylum and genus levels. Chao 1 (Bacterial richness index) and Shannon (Bacterial diversity index) alpha diversity indices were analyzed using QIIME (Version 1.7.0) [28].

2.9. Challenge Test

After 28 days of feeding trial, a challenge test was performed in each group with *A. veronii* which isolated from intestines of diseased crucian carp and preserved in our laboratory. *A. veronii* was inoculated into BHI medium and cultured overnight at 37 $^{\circ}$ C and settled by centrifugation then resuspended and adjusted to sterile phosphate-buffered saline (PBS). Thirty crucian carp were randomly selected from each group and challenged with 0.1 mL of *A. veronii* bacterial suspension (3.6×10^7 CFU/mL) by intraperitoneal injection. Mortality was recorded every 24 h for 10 days. The presence of *A. veronii* as the only etiological agent was confirmed by swabbing the intestinal tissues onto BHI plates and subsequent bacteria identification.

2.10. Statistical Analyses

Statistical analysis was performed using SPSS 19.0 (IBM Corp., Armonk, NY, USA). Differences between groups were detected using one-way analysis of variance (ANOVA) and tested by least significant difference (LSD) test. The results are presented as the mean \pm standard deviation (SD). Differences were considered statistically significant at $p < 0.05$.

3. Results

3.1. Growth Performance

After the feeding trial lasting 28 days, the survival rates of the four crucian carp groups were >95% (Table 2), and there was no significant difference between the groups. Compared with the CG, the fish in EF1 and EF2 groups had a significantly higher final weight, WGR and SCR ($p < 0.05$). Furthermore, SGR of fish in EF3 was significantly higher than in CG group ($p < 0.05$), but there was no difference in final weight or WGR between CG, EF1, and EF2 groups ($p > 0.05$). CF was unaffected by dietary treatment in all groups.

Table 2. Growth performance of crucian carp fed with different experimental diets for 28 days.

Parameters	CG	EF1	EF2	EF3
Initial weight (g)	47.67 ± 0.62	48.60 ± 0.15	48.37 ± 0.52	48.01 ± 0.51
Weight at sampling (g)	64.57 ± 0.93 a	68.44 ± 1.05 b	69.45 ± 0.96 b	67.38 ± 2.74 ab
WGR (%)	35.47 ± 1.11 a	40.83 ± 1.78 b	43.59 ± 0.58 b	40.31 ± 4.26 ab
SGR (%/d)	1.08 ± 0.03 a	1.22 ± 0.04 b	1.29 ± 0.02 b	1.21 ± 0.11 b
Survival rate (%)	96.00 ± 2.00	96.67 ± 2.31	98.00 ± 2.00	96.67 ± 3.01
(CF) (g/cm ³)	2.29 ± 0.16	2.45 ± 0.49	2.41 ± 0.13	2.21 ± 0.18

Note: Each value represents mean ± SD ($n = 3$), and bars with different letters indicate a significant difference by LSD test for each time point ($p < 0.05$). CG, control group; EF1, *E. faecalis* at 10^5 CFU/g diet; EF2, *E. faecalis* at 10^6 CFU/g diet; EF3, *E. faecalis* at 10^7 CFU/g diet.

3.2. Serum Biochemical Parameters

Dietary *E. faecalis* supplementation had a significant effect on some serum biochemical indices of crucian carp (Figure 1). C3 and IgM in the EF2 group increased with the progression of feeding trial and reached maximum values at 2.1 and 10.4 g/L at the end of 28 days of experiment, respectively, which were significantly higher than those in CG and EF1 groups ($p < 0.05$). AKP in the EF2 group was significantly higher than that in CG and EF3 groups at 21 days ($p < 0.05$), and it was significantly higher than in the other three groups at 28 days ($p < 0.05$). There were no significant differences in TG, TCHO, and AST among all groups ($p > 0.05$).

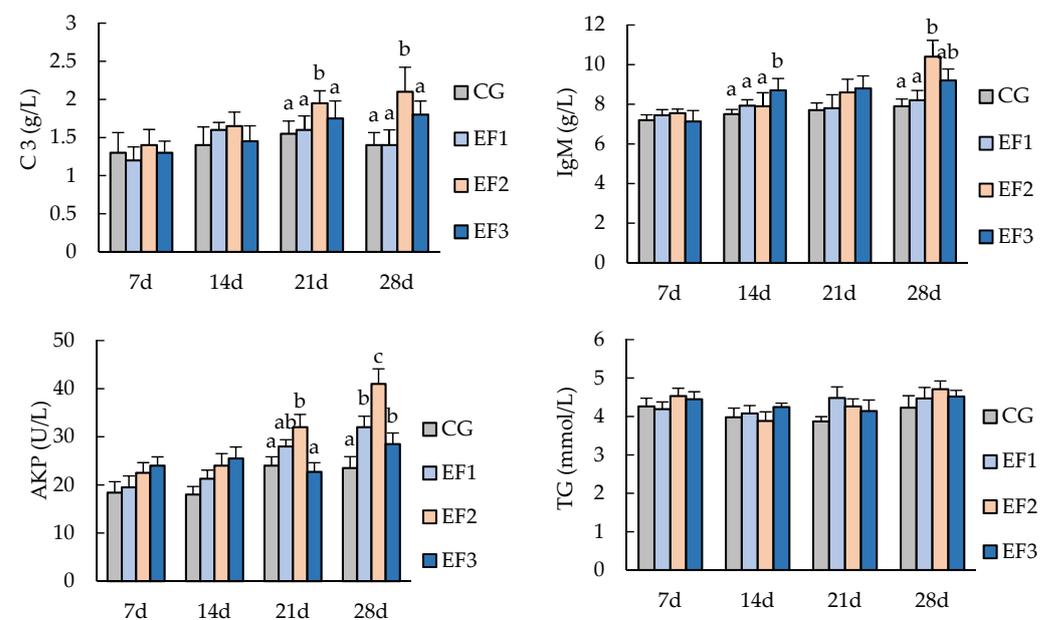


Figure 1. Cont.

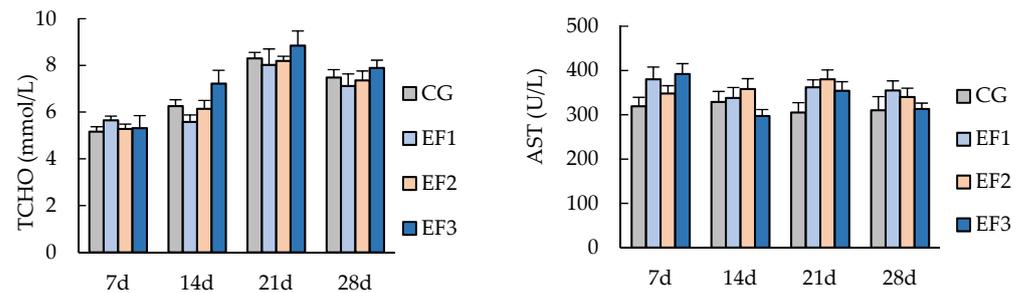


Figure 1. Serum biochemical parameters of crucian carp. Each value represents mean \pm SD ($n = 3$), and bars with different letters indicate significantly differences by LSD test for each time point ($p < 0.05$). C3, complement 3; IgM, immunoglobulin M; TG, triglyceride; TCHO, total cholesterol; AKP, alkaline phosphatase; AST, aspartate aminotransferase; CG, control group; EF1, *E. faecalis* at 10^5 CFU/g diet; EF2, *E. faecalis* at 10^6 CFU/g diet; EF3, *E. faecalis* at 10^7 CFU/g diet.

3.3. Changes in Immunity-Related Genes Expressions

Expression levels of immunity-related genes from the spleen tissue of crucian carp were measured on day 7, 14, 21, and 28, including the genes encoding IL-1 β , C3, and IgM (Figure 2). After 21 days of experiment, the relative expression of IL-1 β in the EF2 group was significantly upregulated compared with CG and EF1 groups ($p < 0.05$), and those in EF1 and EF2 groups were significantly upregulated compared with CG group and EF3 group at 28 days ($p < 0.05$). Expression of C3 in the EF2 group was upregulated at 7 days, and was significantly higher from 14 days than those in other groups ($p < 0.05$). Meanwhile, C3 expression in EF1 was significantly upregulated compared with CG at 21 days and 28 days ($p < 0.05$). Expression of IgM in EF3 group was significantly higher than in CG group from 14 days ($p < 0.05$), and that in EF2 was significantly upregulated from 21 days compared with CG and EF1 groups ($p < 0.05$), with expression peaking at 28 days.

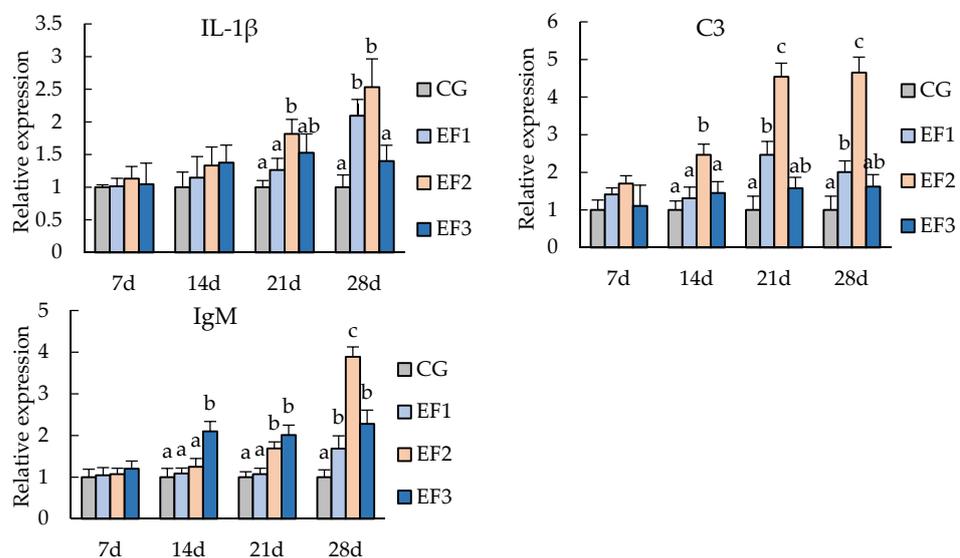


Figure 2. Relative mRNA expression levels of immunity-related genes in crucian carp spleen tissue. Each value represents mean \pm SD ($n = 3$), and bars with different letters indicate significantly differences by LSD test for each time point ($p < 0.05$). CG, control group; EF1, *E. faecalis* at 10^5 CFU/g diet; EF2, *E. faecalis* at 10^6 CFU/g diet; EF3, *E. faecalis* at 10^7 CFU/g diet.

3.4. Changes in Intestinal Morphology

After 28 days of feeding, crucian carp intestines from each fed with *E. faecalis* supplemented diet had intact, orderly, and tight mucosal layers, submucosal layers, muscular layers, and outer membranes (Figure 3). Epithelial cells in the mucosal layer of intestinal tissue in CG group were disordered and vacuolated obviously. In the EF1 group, a few

epithelial cells were ruptured or vacuolated at the top of the mucosal layer. Epithelial cells were intact and arranged normally in EF2 and EF3 groups.

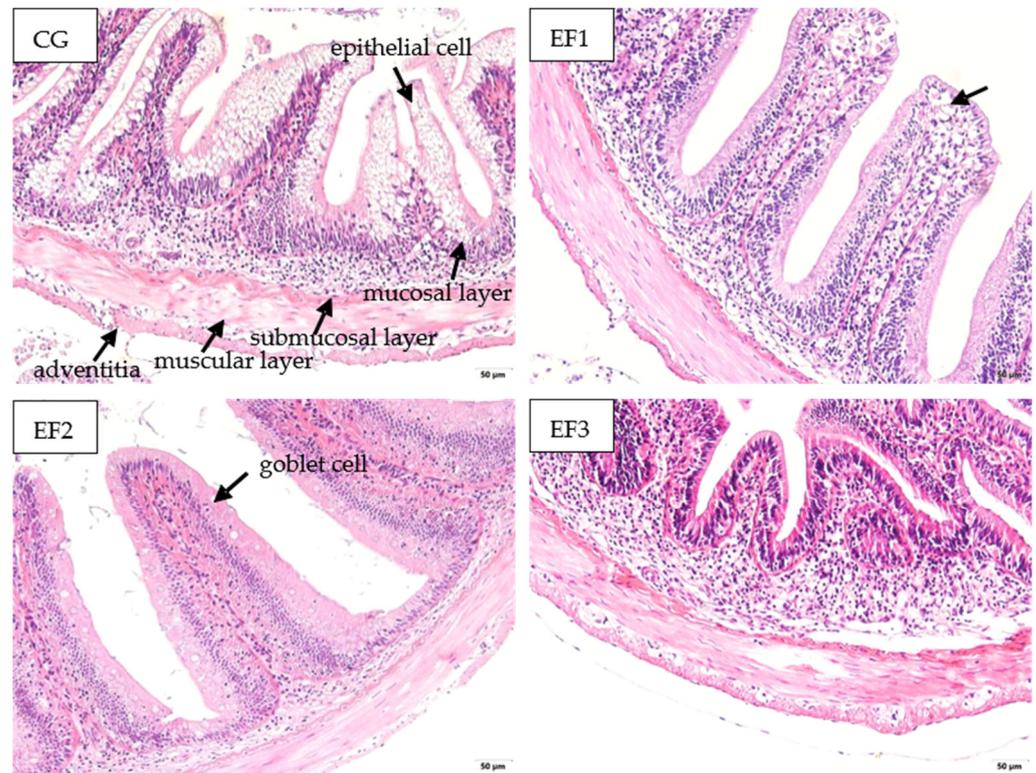


Figure 3. Intestinal morphology analysis of crucian carp fed with different experimental diets for 28 days (200×). CG, control group; EF1, *E. faecalis* at 10^5 CFU/g diet; EF2, *E. faecalis* at 10^6 CFU/g diet; EF3, *E. faecalis* at 10^7 CFU/g diet.

3.5. Changes in the Intestinal Microbiota

We sequenced the 16S rRNA gene amplicons of the intestinal microbiota in the intestine samples of experimental crucian carp, and raw data have been deposited in the NCBI. The average number of OTUs obtained from the different samples ranged from 812 to 1037 (Table 3). The results showed no differences between Chao 1 index and OTUs in GC, EF1, and EF3 groups, but these parameters were significantly lower than in EF2 group ($p < 0.05$). There was no significant difference of Shannon index among all four groups ($p > 0.05$).

To further explore the composition of the microbiota community richness in each group, the abundance in the intestinal microbiota of the top eight bacteria was calculated at the phylum and genus levels. At the phylum level, Proteobacteria, Firmicutes, Actinobacteria, and Bacteroidetes were the primary intestinal microbiota in all groups (Figure 4). Compared with CG group, the relative abundance of Proteobacteria was decreased in EF1, EF2, and EF3 groups, while the relative abundance of Actinobacteria was increased. At the genus level, the primary intestinal microbiota in all groups were *Aeromonas*, *Magnetospirillum*, *Acinetobacter*, and *Cetobacterium* (Figure 4). The relative abundance of *Aeromonas* and *Acinetobacter* was decreased in EF1, EF2, and EF3 groups after 28 days of feeding compared with CG group, while *Cetobacterium* was increased, and EF2 group exhibited the largest differences.

Table 3. OTUs classification and alpha diversity index of microbial communities in the intestine samples.

Sample	OTUs	Chao 1	Shannon
CG	824.67 ± 56.52 a	857.24 ± 46.64 a	5.40 ± 0.31
EF1	842.33 ± 23.51 a	911.79 ± 40.69 a	5.12 ± 0.83
EF2	1025.67 ± 81.24 b	1190.46 ± 102.67 b	5.75 ± 0.09
EF3	878.67 ± 79.74 a	882.85 ± 41.28 a	5.61 ± 0.37

Note: Each value represents mean ± SD ($n = 3$). Different letters indicate significant ($p < 0.05$) difference by LSD test. CG, control group; EF1, *E. faecalis* at 10^5 CFU/g diet; EF2, *E. faecalis* at 10^6 CFU/g diet; EF3, *E. faecalis* at 10^7 CFU/g diet. OTU, operational taxonomic unit.

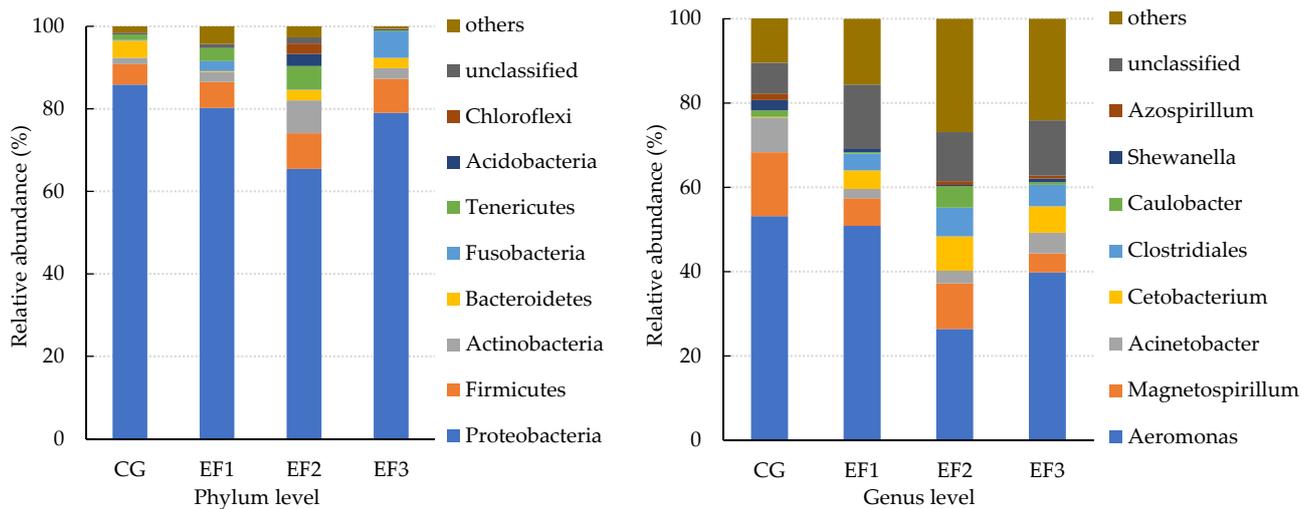


Figure 4. Main relative abundance of the intestinal microbiota in crucian carp at the phylum and genus levels. CG, control group; EF1, *E. faecalis* at 10^5 CFU/g diet; EF2, *E. faecalis* at 10^6 CFU/g diet; EF3, *E. faecalis* at 10^7 CFU/g diet.

3.6. Challenge Test

The cumulative survival rates of crucian carp challenged with *A. veronii* for 10-day are shown in Figure 5. At the end of the 10 days challenge test, the cumulative survival rates were 24.44%, 56.67%, 67.78%, and 61.11% in the CG, EF1, EF2, and EF3 groups, respectively. Crucian carp in treatment groups showed a significantly higher survival rate than those in CG group after 28 days of feeding.

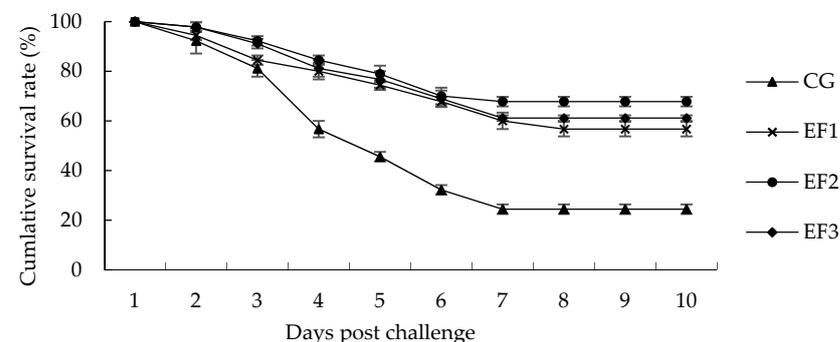


Figure 5. Cumulative survival rate following challenge with *A. veronii* for 10 days. Each value represents the mean ± SD ($n = 3$). CG, control group; EF1, *E. faecalis* at 10^5 CFU/g diet; EF2, *E. faecalis* at 10^6 CFU/g diet; EF3, *E. faecalis* at 10^7 CFU/g diet.

4. Discussion

4.1. Growth Performance

Dietary supplementation of probiotics in aquaculture offers an ecofriendly prophylactic measure [29]. Their applications are increasing due to their positive impact on the

growth and health of aquatic animals [30]. *E. faecalis* as a kind of lactic acid bacterium possessing growth-promoting potential [31]. Previous studies had demonstrated that *E. faecalis* had a positive impact on the growth of tilapia (*Oreochromis niloticus*) [18], rainbow trout (*Oncorhynchus mykiss*) [19], juvenile sea cucumber (*Apostichopus japonicus* selenka) [32], and other aquatic animals [33]. In the present study, following dietary supplementation with *E. faecalis* at 10^5 – 10^7 CFU/g, crucian carp exhibited significantly increased growth after 28 days, consistent with the results of these previous studies.

4.2. Serum Biochemical Parameters

Serum biochemical parameters are the main indices reflecting the health condition and physiological state of fish [34]. Serum parameters related to immunity and antioxidant capacity were measured in this study to confirm the positive effects of the *E. faecalis* on the health status of fish. TG, TCHO, and AST showed no significant differences after dietary intake of *E. faecalis*, in line with similar studies on crucian carp dietary supplementation with *Bacillus subtilis* [35]. C3, the most abundant complement protein in serum, is a bridge between natural and adaptive immunity, and plays an important role in immune defenses, regulation, and pathology [36]. AKP is a lysosomal enzyme involved in macrophage activation, and a promising antimicrobial agent [37]. In this study, C3 and AKP increased with progressing of feeding trial for diets supplemented with *E. faecalis*. Similarly, previous studies showed that dietary *E. faecalis* at 10^6 CFU/g can improve the serum C3 levels and AKP activity in both tilapia and blunt snout bream [18,38]. IgM is the first antibody produced by the body following stimulation by antigen, and it has strong cytotoxic and cytolytic activity [39]. Previous studies revealed higher serum IgM levels in crucian carp fed with *Lactobacillus plantarum* or *Lactobacillus casei* diets [23,40]. In the present study, dietary *E. faecalis* at 10^6 – 10^7 CFU/g may increase serum IgM at different times, similar to the results of these previous studies.

4.3. Immune Related Gene Expressions

Expression of immune-related genes in spleen also promotes the immune responses in fish. Specific immunity that mediate the production of cytokines and antibodies play an important role in the immune responses [24]. IL-1 β plays a central role in the regulation of immune and inflammatory responses to infections by activating lymphocytes or inducing the release of other cytokines [41,42]. IgM, the main agent in humoral immunity, neutralizes antigens and activates the complement cascade [39]. C3 complement plays an antibacterial role by stimulating cell phagocytosis mediated by the complement activation pathway [43]. Heat-killed *E. faecalis* can induce cell-mediated immunity and enhanced IL12 and IFN γ expression in crucian carp [14]. In addition, *E. faecalis* can upregulate the gene expression levels of TNF- α and IL-8 in tilapia [18]. In the present study, crucian carp fed with *E. faecalis* showed significantly increased expression of IL-1 β , C3, and IgM genes in the spleen, consistent with the serum immune indices. This might be related to exopolysaccharides (EPSs), which can promote the expression of immune genes in aquatic animals, since *E. faecalis* is a lactic acid bacterium that synthesizes EPSs [44]. However, the timing at which IL-1 β , C3, and IgM gene expression is first upregulated is inconsistent. This might be related to different pathways of immune gene expression and/or different mechanisms of genes involved in immune regulation [45].

4.4. Intestinal Morphology

In aquatic animals, the intestine is the main organ for food digestion and nutrient absorption [46]. As the largest immune organ in the body, the intestinal tract plays an important role in reducing the invasion of pathogenic bacteria [47]. Studies have shown that dietary ingredients may have an effect on fish intestines, for example, dietary supplementation with *B. cereus* or *B. subtilis* can improve the fold height and microvillus height of the intestines in crucian carp [22,35]. In addition, dietary *E. faecalis* can enhance the intestinal health and nutrient absorption in piglets by increasing the villus height [48]. In

the present study, epithelial cells were intact and arranged normally without ruptured or vacuolation after feeding with *E. faecalis* at 10^6 – 10^7 CFU/g diet. This might be because *E. faecalis* forms a lactic acid barrier on the intestinal mucosa that resists the invasion of pathogenic bacteria and maintains the health of the intestinal tissue [17].

4.5. Changes in the Intestinal Microbiota

The intestinal microbiota plays important roles in intestinal physiology due to its critical impacts on metabolism and immune function [49,50]. Normally, the intestinal microbiota maintains a dynamic balance and stability in the intestinal environment, which can help the body to resist the invasion of pathogenic bacteria and enhance host immunity [51,52]. Diet plays a major role in altering the intestinal microbiota and metabolism [53]. In the present study, the richness of the intestinal microbial community of crucian carp increased after feeding them an appropriate amount of *E. faecalis*. From the perspective of phylum classification, Proteobacteria, Firmicutes, and Actinobacteria were the dominant phyla in all groups. A previous study reported slightly different results [54], possibly due to differences in environmental conditions or feeding habits [55]. Proteobacteria, the largest bacteria phylum, includes many pathogenic bacteria such as *Escherichia coli*, *Salmonella*, and *Vibrio cholerae*, and it has been proposed as a potential signature of dysbiosis and disease risk [56]. Firmicutes are considered beneficial bacteria in the intestines, exerting a positive influence on growth performance, immunity, digestion, and disease resistance in aquatic animals [57]. Actinobacteria can produce secondary metabolites, including extracellular enzymes and potent antibiotics [58]. Our results showed that the abundance of Firmicutes and Actinobacteria was increased after feeding *E. faecalis* for 28 days, but Proteobacteria was decreased compared with the CG group. It is assumed that *E. faecalis* might improve intestinal health by increasing the proliferation of Firmicutes and Actinobacteria, thereby inhibiting the growth of Proteobacteria.

At the genus level, our results showed that *Aeromonas*, *Magnetospirillum*, *Acinetobacter*, and *Cetobacterium* were the dominant genera in all groups. *Aeromonas* and *Acinetobacter* are associated with bacterial fish diseases, and they are also important opportunistic pathogens in crucian carp [5,59]. *Cetobacterium* can be abundant in various freshwater fish species, and it can produce large quantities of vitamin B-12 [60]. In the present study, compared with the control group, the abundance of the *Cetobacterium* was increased in EF1, EF2, and EF3 groups, while *Aeromonas* and *Acinetobacter* were decreased. This might be because *E. faecalis* produces bacteriocins and other bacteriostatic substances that inhibit the growth of pathogenic bacteria [61]. However, in this study, *Enterococcus* did not become the dominant genus. Similarly, the corresponding bacteria were not found in the intestinal of grass carp fed with *Bacillus subtilis* (10^8 CFU/g) for 4 weeks, but the microbiota structure changed [62]. This may be due to the short feeding time, low feeding amount, and other reasons that cannot occupy the intestinal niche, but through the production of some metabolites to regulate the intestinal microbiota structure.

4.6. Disease Resistance

A. veronii is regarded as one of the major pathogens, causing hemorrhagic septicemia, ulcerations, and ascites in crucian carp [63]. *A. veronii* exhibits significant virulence toward crucian carp, with an LD50 value of 1.31×10^7 CFU/mL [5]. In the present study, after a 10-day challenge with 3.6×10^7 CFU/mL *A. veronii*, the cumulative survival rate was 24.44% in the control group. Meanwhile, fish in the three *E. faecalis* groups all showed significantly higher cumulative survival rates, with the EF2 group showing the highest rate (67.78%). Similar results were observed in crucian carp fed diets containing *Bacillus velezensis* and *Lactobacillus casei* supplemented-diets [23,63]. This might be because *E. faecalis* can increase the abundance of intestinal probiotics and stimulate cellular and humoral immune functions, including complement C3 and IgM, to protect hosts against pathogen infection.

5. Conclusions

In this study, the addition of *E. faecalis* YFI-G720 to the diet positively influenced the growth performance, immune response, the intestinal microbiota structures, and resistance to pathogens in crucian carp. Long-term intake of fresh *E. faecalis* YFI-G720 at 10^6 CFU/g achieved a more pronounced effect. Therefore, *E. faecalis* YFI-G720 can be used as a potential probiotic in crucian carp farming. However, the reason that *E. faecalis* YFI-G720 improves the immunity of crucian carp is still unclear, and we will investigate the metabolites and functions of *E. faecalis* YFI-G720 to provide a reference for the efficient use.

Author Contributions: Y.Z., Y.X. and Y.L. conceived and designed the study, and performed the data collection, analysis, statistical analysis, writing of the manuscript, and conducted the software and literature review. Y.X. and Z.X. conducted animal management and sample collections. M.X., Y.L. and Z.X. performed the microbial analysis, immunity analysis, and literature review. Y.X., Y.Z., Y.F. and L.Z. reviewed and edited the manuscript. All authors have read and agreed to the published version of the manuscript.

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Institutional Review Board Statement: In the present study, all experimental procedures were conducted according to guidelines of the appropriate Animal Experimental Ethical Inspection of Laboratory Animal Center, Yangtze River Fisheries Research Institute, Chinese Academy of Fishery Sciences (ID Number: YFI2021-zhouyong-02).

Data Availability Statement: The data presented in this study are available on request from the corresponding author.

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