The Effect of Dietary Saccharoculture on Growth Performance, Non-Specific Immunity and Autochthonous Gut Microbiota of Gibel Carp *Carassius auratus*

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Abstract

The objective of this study was to evaluate the effect of a commercial fermented yeast product (Saccharoculture, Veterinary Pharmaceuticals, Korea), on the growth performance, non-specific immunity and autochthonous intestinal microbiota of gibel carp *Carassius auratus*. Following an eight week feeding trial the growth performance and serum non-specific immunity of fish fed one of four treatments (A: control group, B: 2 g/kg Saccharoculture, C: 3 g/kg Saccharoculture or D: 2 g/kg Saccharoculture + 0.1 g/kg flavomycin) was measured. Additionally, the predominant autochthonous intestinal microbiota was analyzed by denaturing gradient gel electrophoresis (DGGE). The results indicated that dietary supplementation of Saccharoculture or Saccharoculture combined with flavomycin significantly improved final weight, weight gain (WG), special growth rate (SGR) and feed conversion ratio (FCR) compared to the control group (*P* < 0.05), and the best growth performance was obtained in group B. However, the non-specific immunity factors were not significantly affected by dietary yeast or flavomycin (*P* > 0.05). Compared to the control group, fish fed dietary yeast exhibited modulated intestinal bacterial communities. Levels of some bacteria were elevated by dietary Saccharoculture, such as, *Acinetobacter* sp. (FR749840.1, FJ646641.1), *Escherichia vulneris* (HQ259947.1) and *Cetobacterium somerae* (AB353124.1). *Cetobacterium* sp. (HM778168.1) and *Lactococcus lactis* subsp. (EF589778.1) were suppressed in the Saccharoculture with flavomycin group. However, *Nevskia* sp. (AB426559.1) was suppressed by the higher level Saccharoculture or Saccharoculture and flavomycin combination.

Keywords: Fermented yeast product; Gibel carp; Intestinal microbiota; DGGE

Introduction

Dietary supplementation with a variety of products containing yeast (*Saccharomyces cerevisiae*) has been evaluated in a number of different fish species [1-4]. It has been observed that fermented yeast products can positively influence growth performance [1,2,4], as well as the non-specific immune responses, such as lysozyme and complement activity [1,3,4]. The efficacy of yeast maybe contributed to its various immune stimulating compounds such as β-glucans, nucleic acids, mannan oligosaccharides (MOS) and other cell wall components [5].

The gibel carp (*Carassius auratus*) is an important fish species in China, because of its excellent taste and rapid growth [6]. The fish intestinal microflora has been implicated as playing several functions in respect to nutritional digestion, defending against pathogens and enhancing immunity [7-9]. However, there is no information on the effects of dietary yeast on the intestinal microbiota of gibel carp. Also in China, antibiotic growth promoters such as flavomycin are widely used to improve production to satisfy the growing demand for aquaculture products [4]. To the best of our knowledge, no information is available on the effects of yeast supplementation or yeast combined with flavomycin on gibel carp. Thus the present study was conducted to compare the effect of dietary Saccharoculture and a Saccharoculture and flavomycin combination on growth parameters, non-specific immunity and intestinal microbiota of gibel carp.

Methods

Experimental diets

The basal diet formulation and proximate composition are shown in Table 1; the chemical composition was analyzed according to AOAC methods [10]. The basal formulation served as the control diet and three experimental diets were produced by supplementing the basal formulation with yeast and flavomycin, as shown in Table 2. The yeast product, Saccharoculture, was supplied by Veterinary Pharmaceuticals, Korea; the level of probiotic *Saccharomyces cerevisiae* J8734 is ca. 1.5×10⁹ CFU per kilogram of Saccharoculture. The diets were extruded to obtain pellets (2-3 mm diameter), then were dried using an electrical fan at room temperature and stored at 4°C until feeding.

Husbandry conditions

Gibel carp were obtained from a fish farm in Hefei city, Anhui, P.R. China. Fish were acclimated to laboratory conditions for 15 days in a 1000 L plastic tank and fed a basal diet twice a day during the period. The feeding trial was carried out in a recirculation system consisting of 20 flat bottom fiberglass tanks (60 cm × 60 cm × 60 cm, effective water volume 180 L). Tanks were connected to a central processing system.

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Randomly sampled to measure body length, body height, body width, satiation twice a day (at 9:00 and 16:00). One day, and 460 fish were selected, batch-weighed and allocated into by fluorescent lamps from 8:30 to 20:00. Dissolved oxygen was maintained above 6.00 mg/L, ammonia below ammonia. Aeration was provided intermittently for 3 min per 15 min. Clinoptilolite is a natural zeolite with a strong adsorption capacity for with column clinoptilolite filter (volume, 1500 L) for water treatment.

\[
\text{Slaughter index (SI)} = 100 \times \frac{\text{body weight without viscera (g)}}{\text{body weight (g)}}
\]

\[
\text{Viscerosomatic index (VSI)} = 100 \times \frac{\text{viscera weight (g)}}{\text{body weight (g)}}
\]

Where IBW is the initial body weight of fish, FBW is the final body weight of fish, t is the experimental period (56 days) and IT\text{a} is the total dry diet consumption during the experimental period.

**Non-specific immunological analysis**

After the termination of the feeding trial, three fish from each tank were sampled for blood. Blood was taken from caudal vein by hypodermic syringe, and blood was left at 4°C overnight, then centrifuged at 3000 g for 10 min, and isolated sera was stored frozen at -20°C for subsequent analysis of lysozyme (LSZ), complement 3 (C3) and complement 4 (C4) activity. Serum LSZ activity, the activity of serum C3 and C4 were measured according to He et al. [4].

**Intestinal predominant microbiota analysis**

Five fish from each tank were randomly sampled at the end of the feeding trial. The intestinal tracts were sampled and processed as described elsewhere [11]. To avoid individual variation, the sampled intestinal walls from each tank were homogeneously mixed together [9,11,12]. The total genomic DNA was extracted with CTAB and lysozyme methods [4] and purified with a Tiangen DNA purification kit (Tiangen Biotechnologies, Beijing, China).

The V3 region of the 16S rRNA gene was amplified according to Liu et al. [12]. DGGE of PCR products was performed with the Bio-Rad Dcode mutation detection system (Bio-Rad, Hercules, CA, USA) according to [4,11,12]. To identify the inserted sequences, the BLAST 2.0 algorithm was used to compare the derived sequence to 16S rRNA sequences in the DNA Data Bank of Japan (DDBJ) database. Species identification was made on the basis of percentage similarity to database sequences (98.0–100.0% similarity).

**Statistical analysis**

Data are expressed as mean values ± S.D. The effects of diet on immune parameters and growth performance were analyzed by one-way analysis of variance (ANOVA) using statistical analysis software (SAS) (Cary, NC, USA). In our study, a similarity coefficient (Cs) matrix less than 0.60 is regarded as significant difference; while 0.60 < Cs < 0.85 is marginal difference and Cs > 0.85 is very similar [4].

**Results**

**Growth performance**

The growth performance of carp after the 56 d experiment is shown in Table 3. The weight gain (WG) and final body weight (FBW) of carp in groups B (2 g/kg Saccharoculture) and D (2 g/kg Saccharoculture + 0.1 g/kg flavomycin) were significantly higher than the control group and group C (P < 0.05); the highest level appeared in group B. The higher level of dietary Saccharoculture (Group C) did not significantly affect WG or FBW compared to the control group (P > 0.05). Specific growth rate (SGR) was significantly affected by the dietary fermented yeast product (Table 3, P < 0.05), except for group C. The highest SGR was obtained in group B and then D, both of which were significantly higher than the control group and group C (P < 0.05).

The condition factor (BI) of fish in group B and D were significantly

**Table 1:** Feed formulation and chemical composition of the basal diet used in the experiment.

| Diet       | Basal diet | Saccharoculture | Flavomycin B% |
|------------|------------|-----------------|--------------|
| A          | 100        |                 |              |
| B          | 99.8       | 0.2             |              |
| C          | 99.7       | 0.3             |              |
| D          | 99.7       | 0.2             | 0.01         |

1 supplied by Veterinary Pharmaceuticals, Korea
2 supplied by Xiangweisi Ltd, Shangdong, China

with column clinoptilolite filter (volume, 1500 L) for water treatment. Clinoptilolite is a natural zeolite with a strong adsorption capacity for ammonia. Aeration was provided intermittently for 3 min per 15 min. Dissolved oxygen was maintained above 60.0 mg/L, ammonia below 0.20 mg/L and temperature at 25±0.5°C. The illumination was provided by fluorescent lamps from 8:30 to 20:00.

**Experimental design**

Prior to the beginning of the experiment, the fish were starved for one day, and 460 fish were selected, batch-weighed and allocated into each of the twenty tanks (23 fish per tank). Each experimental diet had four replicate tanks. During the experiment, the fish were fed to satiation twice a day (at 9:00 and 16:00).

**Growth measurements**

The trial lasted for 56 days. At the end of the trial, fish were weighed after one day of food deprivation, and five fish from each tank were randomly sampled to measure body length, body height, body width, viscera and hepatopancreas weights. Growth performance, feed utilization, viscerosomatic index and slaughter index were calculated according to the following formulae:

Weight gain (WG) = \(100 \times (\text{FBW} - \text{IBW}) / \text{IBW}\);

Special growth rate (SGR) = \(100 \times \ln(\text{FBW} / \text{IBW}) / t\);

Feed conversion rate (FCR) = IT\text{a} (g) / WG (g);

Condition factor (BI) = bodyweight (mg) / \([\text{body length (cm)} \times \text{body height (cm)} \times \text{body width (cm)}]\);
IBW, initial body weight; FBW, final body weight; WG, weight gain; SGR, specific growth rate; different letters showed significant differences (P < 0.05).

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| Parameters   | A        | B          | C          | D          |
|--------------|----------|------------|------------|------------|
| IBW(g)       | 7.55±0.06 | 7.54±0.06  | 7.55±0.13  | 7.56±0.15  |
| FBW(g)       | 28.12±1.08 | 33.55±0.71 | 29.08±0.39 | 32.29±1.04 |
| WG(%)        | 27.14±14.95 | 34.4±11.34 | 25.44±17.77 | 32.34±17.00 |
| SGR(%)       | 2.34±0.07  | 2.66±0.05  | 2.38±0.07  | 2.58±0.07  |
| FCR          | 1.62±0.07  | 1.49±0.05  | 1.53±0.05  | 1.50±0.06  |

* Superscripts showed the result of multiple range test (Duncan’s procedure). Different letters showed significant differences (P < 0.05).

Table 3: Effects of experimental diets on the growth and diet utilization of gibel carp Carassius auratus.

| Diet          | BI (mg/cm3) | SI (%) | VSI (%) |
|---------------|-------------|--------|---------|
| A             | 456.3±12.51 | 89.02±1.55 | 4.22±0.15 |
| B             | 483.2±11.94 | 89.10±1.44 | 3.98±0.22 |
| C             | 445.8±17.45 | 90.15±1.05 | 3.74±0.32 |
| D             | 479.6±10.29 | 89.50±1.27 | 3.76±0.28 |

* Superscripts showed the result of multiple range test (Duncan’s procedure). Different letters showed significant differences (P < 0.05).

Table 4: Effects of experimental diets on the condition factor (BI), slaughter index (SI) and viscerosomatic index (VSI) of gibel carp Carassius auratus.

** Superscripts showed the result of multiple range test (Duncan’s procedure). Different letters showed significant differences (P < 0.05).

Table 5: Effects of experimental diets on the serum lysozyme (LYZ) and complement activity (C3 and C4) of gibel carp Carassius auratus.

Discusstion

In the present study, dietary supplementation with yeast had positive effects on gibel carp growth rate, weight gain and SGR compared to the control diet (Table 4). Similar benefits have been reported previously in fish fed dietary yeast [13,14]. The beneficial influence of Saccharoculture on growth was possibly due to its high content of nucleic acids, mannan oligosaccharides (MOS), lactose and other cell wall components [15]. Previously it has been reported that Japanese flounder fed a diet supplemented with 5 g kg−1 MOS exhibited better weight gain and feed conversion ratio than those fed a control diet [16]. However, in the present study, the growth performance of fish fed 0.3% Saccharoculture was lower compared to that of fish fed 0.2% Saccharoculture (Table 4). Several reports have suggested that low level dietary inclusion of yeast products could efficiently improve aquatic animal growth parameters; 0.1% inclusion of brewer’s yeast has been reported to elevate the growth performance and feed utilization of Nile tilapia after 9 weeks feeding [2] and 0.25% baker’s yeast has been reported to improve gold fish growth parameters [17]. But even higher levels of dietary commercial brewer’s yeast product (2%) could also exert a positive influence on the growth performance of juvenile beluga [18]. These contradictory results may be attributed to the different fish species and different properties of the yeast products. In the present study, the growth performance was not further strengthened by dietary inclusion of the antibiotic growth promoter flavomycin.

Yeast cells provide about 7.7% crude glucan [19], and glucans have...
been reported to be capable of enhancing innate immune responses, including respiratory burst of head kidney macrophages, serum complement activity and lysozyme activity [20-22] when administered by injection. However, the increase of serum lysozyme was not observed in fish orally receiving glucans [23], which agrees with the results of the present study. The complement system is an important

| phylum          | Band No. | Closest relative (obtained from BLAST search) | Identity (%) | A       | B       | C       | D       |
|-----------------|----------|-----------------------------------------------|--------------|---------|---------|---------|---------|
| Proteobacteria  | 1        | *Acinetobacter* sp. (HQ841068.1)               | 100          | 0.8±1.1 | 0.0     | 0.0     | 0.9±0.2 |
|                 | 2        | *Acinetobacter* sp. (EU260174.1)               | 99           | 2.2±0.5 | 1.1±0.6 | 0.9±0.7 | 1.4±0.9 |
|                 | 4        | *Acinetobacter* haemolyticus (HQ132734.1)     | 100          | 1.3±0.3 | 0.9±0.2 | 0.5±0.2 | 0.8±0.5 |
|                 | 5        | *Acinetobacter* sp. (FR749840.1)              | 100          | 19.4±7  | 23.0±1.7| 25.6±3.2| 12.0±7.3|
|                 | 6        | *Acinetobacter* sp. (HQ59186.1)               | 100          | 1.2±0.6 | 0.7±0.0 | 0.8±0.2 | 0.9±0.3 |
|                 | 9        | *Acinetobacter* sp. (FJ364641.1)              | 99           | 1.2±0.4 | 1.9±0.2 | 1.5±0.6 | 0.8±0.4 |
|                 | 13       | *Nevskia* sp. (AB426558.1)                    | 100          | 4.3±1.2 | 4.2±1.0 | 1.4±1.0 | 2.8±1.5 |
|                 | 14       | *Escherichia* vulneris (HQ259947.1)           | 100          | 0.5±0.2 | 1.4±0.8 | 0.7±0.3 | 0.3±0.1 |
|                 | 15       | *Shewanella* xiamenensis (HQ411942.1)         | 100          | 0.0     | 0.0     | 0.0     | 2.6±1.3 |
|                 | 21       | *Escherichia* fergusonii (HQ259962.1)         | 100          | 0.4±0.3 | 1.6±0.8 | 0.2±0.2 | 0.3±0.1 |
|                 | 22       | *Cronobacter* dublinensis (FJ880412.1)        | 100          | 0.0     | 0.7±0.1 | 0.2±0.3 | 0.0      |
|                 | 25       | *Serratia* sp. (HQ588852.1)                   | 100          | 2.6±2.7 | 3.0±0.7 | 1.9±0.6 | 2.7±0.1 |
| Firmicutes      | 10       | *Lactococcus* lactis subsp. lactis (EF589778.1)| 100          | 1.6±0.9 | 0.7±0.5 | 0.6±0.2 | 1.3±0.4 |
|                 | 16       | *Staphylococcus* pasteurii (HQ739095.1)       | 100          | 2.4±2.5 | 0.6±0.1 | 0.4±0.3 | 2.5±0.0 |
|                 | 17       | *Anoxybacillus* sp. (HQ696615.1)              | 100          | 0.7±0.1 | 0.9±1.0 | 0.3±0.2 | 0.4±0.3 |
|                 | 19       | *Lactococcus* lactis subsp. (EF589778.1)      | 99           | 13.8±3.5| 17.1±2.1| 11.6±3.8| 8.7±5.2 |
|                 | 24       | *Anoxybacillus* sp. (FN432807.1)              | 100          | 0.9±1.2 | 0.0     | 0.0     | 1.0±0.2 |
|                 | 26       | *Verrucomicrobiaceae* (HM142713.1)            | 100          | 0.0     | 0.0     | 0.8±0.5 | 0.0      |
|                 | 27       | *Lactococcus* lactis subsp. cremoris (JF297369.1)| 99           | 0.9±0.6 | 0.0     | 2.4±1.0 | 0.7±0.1 |
| Fusobacteria    | 11       | *Cetobacterium* sp. (HM778168.1)              | 100          | 3.2±0.3 | 1.1±0.3 | 4.8±2.1 | 1.7±1.5 |
|                 | 18       | *Cetobacterium* somerae (AB351214.1)          | 100          | 1.9±0.4 | 2.4±0.4 | 3.7±1.3 | 1.2±0.7 |
| Unclassified    | 7        | Marine sponge bacterium (EU346455.1)          | 99           | 0.3±0.3 | 0.0     | 0.0     | 0.3±0.0 |
| bacteria        | 8        | Uncultured bacterium (GU485261.1)             | 99           | 39.6±5.6| 35.7±1.5| 37.6±3.8| 22.6±17.0|
|                 | 12       | Uncultured bacterium clone 2701 (HM452218.1)  | 99           | 1.6±0.7 | 2.2±0.3 | 3.5±0.9 | 1.1±0.5 |
|                 | 20       | Activated sludge bacterium (GU136512.1)       | 99           | 0.0     | 0.2±0.2 | 0.0     | 0.0      |
|                 | 23       | Bacterium B12011 (HQ674993.1)                 | 99           | 0.0     | 0.4±0.1 | 0.3±0.1 | 0.0      |

Table 6: Identity of the gut autochthonous bacterial community of gibel carp *Carassius auratus* fed experimental diets and the relative abundance (%)
element of both the innate and adaptive immune system in fish [24-
26]. In our study, the C3 activity was slightly improved by dietary yeast supplemen-
tation, but this was not significant at \( P < 0.05 \). These effects have been described by Ra" [27] who investigated immunostimulants in both fish and shellfish. The results obtained from recent trout and carp trials show that the serum complement activity of fish fed MOS increases during the investigation period [28].

On comparing the DGGE band patterns in Figure 1, we observed the increasing levels of some bacterium by dietary yeast [Table 6]; these include Acinetobacter sp. (FR749840.1), Lactococcus lactis subsp. (EF89778.1), Cetobacterium somerase (AB353124.1), uncultured bacterium clone 2701 (HM452218.1) and Bacterium B81(2011) (HQ674993.1). Acinetobacter sp. is a normal intestinal bacteria of some fish [4,29], in our lab, two species of Acinetobacter were isolated from the gut of hybrid tilapia (\( C \) O. niloticus \( C \) O. aureas), which could degrade N-Acyl homoserine lactones (AHL) molecules (quorum sensing molecules) produced by Aeromonas hydrophilia (data not published). Acinetobacter sp. have also been reported to degrade AHLS from phytopathogenic bacterium Burkholderia [30]. Therefore elevated gut Acinetobacter sp. may provide a level of protection against Gram negative pathogens, but further studies are required to verify this hypothesis. From our sequencing results, several lactic acid bacteria (LAB) were detected in the intestine of carp. LAB have been considered beneficial residents of the fish's intestinal ecosystem by producing bacteriocins and lactic acid, which inhibit growth of certain fish pathogens and thus positively affect the host's microflora [31-33]. In the present study, DGGE based analysis of the autochthonous intestinal microbiota revealed that L. lactis subsp. (EF89778.1) levels were significantly higher in carp fed 0.2% dietary Saccharoculture (\( P < 0.05 \)); a similar result has been reported with traditional culture-based method [34]. He et al. [4] showed that a Cetobacterium someras-like organism was selectively stimulated in the intestine of hybrid tilapia by dietary yeast culture (DVAQUA, USA), which was also observed in our study. Cetobacterium somerae has been reported to be an important intestinal bacterium in fish which contributes towards the host's vitamin B12 requirements [35]. The results of the present study indicate that the Saccharoculture modulates the intestinal microbiota towards a potentially more beneficial microbial community. However it should be noted that the level of two bacteria, Acinetobacter sp. (FR749840.1) and L. lactis subsp. (EF89778.1), were dramatically reduced with the dietary inclusion of flavomycin. The reason may be that flavomycin have also been reported to degrade flavomycin mainly impaired the transglycosylase activities through penicillin-

binding proteins [36]. Further studies are required to identify the LAB species or other probiotics affected by dietary yeast in order to ascertain their contributory affect towards the host benefits.

In summary, the results indicate that a diet containing 0.2% Saccharoculture with or without 0.01% flavomycin (8%) had the most significant growth-promoting effects on gibel carp. Saccharoculture increased the relative abundance of some potentially beneficial bacteria, such as, Lactococcus lactis subsp. lactis (EF89778.1). However, when Saccharoculture and flavomycin were combined together, flavomycin overshadowed the effect of Saccharoculture on the intestinal microbiota. None of the diets tested exerted effects on the immune parameters tested after 8 weeks.

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