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Dietary α -ketoglutarate alleviates glycinin and β -conglycinin induced damage in the intestine of mirror carp (*Cyprinus carpio*)

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This study investigated the glycinin and β -conglycinin induced intestinal damage and α -ketoglutarate alleviating the damage of glycinin and β -conglycinin in intestine. Carp were randomly divided into six dietary groups: containing fish meal (FM) as the protein source, soybean meal (SM), glycinin (FMG), β conglycinin (FMc), glycinin+1.0% α -ketoglutarate (AKG) (FMGA), β -conglycinin +1.0% AKG (FMcA). The intestines were collected on 7th, and the hepatopancreas and intestines were collected on 56th. Fish treated with SM and FMc displayed reduced weight gain, specific growth rate, and protein efficiency. On 56th day, Fish fed on SM, FMG and FMc presented lower superoxide dismutase (SOD) activities. FMGA and FMcA had higher SOD activity than those fed on the FMG and FMc, respectively. In intestine, fish fed on the SM diets collected on 7th presented upregulated the expression of transforming growth factor beta (TGF β 1), AMP-activated protein kinase beta (AMPK β), AMPK γ , and acetyl-CoA carboxylase (ACC). Fish fed FMG presented upregulated expression of tumor necrosis factor alpha (TNF- α), caspase9, and AMPK γ , while downregulated the expression of claudin7 and AMPK α . FMc group presented upregulated expression of TGF β 1, caspase3, caspase8, and ACC. Fish fed FMGA showed upregulated expression of TGFβ1, claudin3c, claudin7, while downregulating the expression of TNF- α and AMPKy when compared to fish fed FMG diet. FMcA upregulated the expression of TGF β 1, claudin3c than fed on the FMc. In intestine, the villus height and mucosal thickness of the proximal intestine (PI) and the distal intestine (DI) were decreased and crypt depth of the PI and mid intestine (MI) were increased in SM, FMG and FMc. In addition, fish fed on SM, FMG and FMc presented lower citrate synthase (CS), isocitrate dehydrogenase (ICD), α ketoglutarate dehydrogenase complex (α -KGDHC) Na⁺/K⁺-ATPase activity in DI. FMGA had higher CS, ICD, α -KGDHC, and Na⁺/K⁺-ATPase activity in PI and MI than those fed on the FMG. FMcA had higher Na^+/K^+ -ATPase activity in MI. In conclusion, dietary soybean meal destroys the intestine's health, the adverse

effects are related to the presence of β -conglycinin and glycinin, especially glycinin. AKG may regulate intestinal energy *via* tricarboxylic acid cycle, thereby alleviating the damage intestinal morphology caused by the dietary soybean antigen proteins.

KEYWORDS

α-ketoglutarate, glycinin, β-conglycinin, intestine immunity, Cyprinus carpio

1 Introduction

Given the increased demand and limited capture of forage fish, fishmeal prices have reached all-time high, with limited supply (1). As an alternative to fishmeal, soybean meal is a plant-based ingredient with high levels of protein, balanced amino acid profile, and other nutrients required by aquatic animals (2). However, soybeanmeal cannot completely replace fishmeal because of the antinutritional factors such as allergens, trypsin inhibitor and saponins (3). High inclusion of soybean meal in fish aquafeeds decreased growth performance in Japanese seabass (Lateolabrax japonicus) (4), orange-spotted grouper, and (Epinephelus coioides) (5). These adverse effects are often related to the presence of anti-nutritional factors, especially β -conglycinin and glycinin (6). These two antigenic proteins are considered the major cause of reduced growth and increased intestinal inflammation in aquatic animals (7). Previous studies have showed that 4% to 8% β-conglycinin and/or glycinin can damage the structural integrity of the intestine, reduce the immune function and compromise the growth performance of juvenile golden crucian carp (Carassius auratus) (8, 9) and grass carp (Ctenopharyngodon idella) (10, 11).

In this context, supplementation of functional additives is an important strategy to maintain the intestine health. α -ketoglutaric acid (AKG) is an intermediate for the tricarboxylic acid (TCA) cycle is involved in pleiotropic metabolic and regulatory pathways in the cell, including energy production (12). AKG has the function of improve intestinal immunity and inhibit apoptosis in response to inflammatory stimuli (13), which can ultimately compensate the energy consumption in the intestinal health. In addition, AKG has alleviative effect on oxidative stress as a source of energy and an antioxidant in mammalian cells (14). Similar results were also observed on juvenile red drum (*Sciaenops ocellatus*) (15). Thus, it is hypothesized that AKG could protect fish against glycinin and β -conglycinin injury.

Nevertheless, the alleviating effects of α -ketoglutarate on the damage of glycinin and β -conglycinin in intestinal mucosa of Mirror carp have not been elucidated. Our research group showed that the intestinal inflammation and tight junction protein showed a trend of first serious and then recovery when addition β -conglycinin and Glycinin was 4% to 16%, respectively (16). Therefore, the objective of this study was to investigate the effects of high-lever glycinin and β -conglycinin on growth

performance, antioxidant capacity, intestinal health and energy of mirror carp (*Cyprinus carpio*), and evaluated whether AKG supplementation had potential alleviating effects on the glycine and β -conglycine induced injuries in the intestinal mucosa.

2 Materials and methods

2.1 Experimental diets

Glycinin (purity, 85%) and β -conglycinin (purity, 88%) were provided by the China Agricultural University. α-ketoglutarate (purity, 98%) was purchased from Sigma-Aldrich. For this study, six experimental diets were formulated to be isonitrogenous (33%) and isolipidic (6%). The ingredients and nutrient composition of all diets are shown in Table 1. For the control group (FM), fish meal was used as the main protein source, and the other five diets were formulated with either 56.0% soybean meal (SM), 22.8% glycinin (FMG), 22.0% β-conglycinin (FMc), 22.8% glycinin + 1.0% αketoglutarate (FMGA), or 22.0% β -conglycinin + 1.0% α ketoglutarate (FMcA). Dietary ingredients were ground into fines to pass through a 60 mesh sieve. Homogenized ingredients were thoroughly mixed with fish oil, soybean oil, phospholipid, and water, and mixtures were extruded (twin screw extruded F-26, South China University of Technology, Guangzhou, China), using a 1.5 mm die-plate. The diets were dried in a ventilated oven at 40 ° C, until reached a constant dry matter, and stored at -20 °C until feeding.

2.2 Feeding management

The feeding trial was carried out at the College of Life Science, Huzhou University. Prior commencement of the feeding trial, fish were acclimated in the recirculating system for two weeks. A total of 450 carps of similar sizes (mean initial body weight 2.08 ± 0.06 g) were randomly distributed into six treatments, 3 replicate tanks per treatment, with 25 fish per tank (500 L water). Fish were fed to apparent satiation thrice daily (08:00, 13:00, and 18:00) for 8 weeks; and uneaten feed was collected after 45 min, dried and weighed to record feed intake (17). During the experimental period, the carp were cultured in a recirculating aquaculture system with continuous aeration, with 1/3 of the water being changed every day. The water TABLE 1 Formulation of the experimental diets fed to Mirror carp (Cyprinus carpio), and their analyzed nutrient composition.

In our disconte	Groups						
Ingredients	FM	SM	FMG	FMc	FMGA	FMcA	
Fish meal	38.00	5.00	5.00	5.00	5.00	5.00	
Soybean meal	0.00	56.00	0.00	0.00	0.00	0.00	
Glycinin	0.00	0.00	22.80	0.00	22.80	0.00	
β-conglycinin	0.00	0.00	0.00	22.00	0.00	22.00	
α-ketoglutarate	0.00	0.00	0.00	0.00	1.00	1.00	
Wheat middling	49.90	28.72	57.10	57.80	56.10	56.80	
Cellulose	3.60	0.00	3.65	3.69	3.65	3.69	
Fish oil	0.00	3.20	3.20	3.20	3.20	3.20	
Soybean oil	3.10	0.30	1.40	1.40	1.40	1.40	
Soybean lecithin oil	1.00	1.00	1.00	1.00	1.00	1.00	
Vitamin premix ^a	0.50	0.50	0.50	0.50	0.50	0.50	
Mineral premix ^b	0.20	0.20	0.20	0.20	0.20	0.20	
Choline chloride	0.30	0.30	0.30	0.30	0.30	0.30	
Ca(H ₂ PO ₄) ₂	1.20	2.00	2.00	2.00	2.00	2.00	
Carboxymethyl cellulose	2.00	2.00	2.00	2.00	2.00	2.00	
Lysine	0.10	0.23	0.00	0.00	0.00	0.00	
Methionine	0.10	0.51	0.65	0.65	0.65	0.65	
L-Threonine	0.00	0.04	0.20	0.26	0.20	0.26	
Proximate analysis (%)							
Moisture	5.70	5.90	5.87	6.30	5.88	5.25	
Crude protein	33.36	34.08	33.62	33.42	33.59	33.75	
Crude lipid	6.50	6.35	6.48	6.42	6.40	6.34	
Ash	11.52	7.41	4.54	4.73	4.66	4.61	

^aThe premix provided the following per kg of diets: A 8 000IU, VC 500mg, VD 3 000IU, VE 60mg, VK 35mg, VB₁ 15mg, VB₂ 30mg, VB₆ 15mg, VB₁₂ 0.5mg; ^bThe premix provided the following per kg of diets:FeSO₄•H₂O 960 mg, CuSO₄•5H₂O 12 mg, MnSO₄•H₂O 33 mg, ZnSO₄•H₂O 70 mg, Na₂SeO₃ 1.2 mg, Ca(IO₃)₂ 1.4mg, CoCl₂•6H₂O 2.4 mg, Zeolite meal 920 mg.

temperature ranged from 24.6 to 30.4 °C, dissolved oxygen was no less than 6.0 mg l^{-1} , and the ammonia nitrogen concentrations was less than 0.3 mg L^{-1} .

Survival rate (SR, %) = $N_t/N_0 \times 100$;

Protein efficiency (PE, %) = $(FBW - IBW)/(F \times P) \times 100;$

Feed conversion ratio (FCR) = F/(FBW - IBW);

Abbreviations: Number of initial fish, N_0 ; number of final fish, N_t ; initial body weight, IBW; final body weight, FBW; Feeding days, d; total diet intake, F; feed crude protein content, P.

2.4 Sample collection and analysis

Intestine samples collected on the 7th day were dissected quickly on ice. 3 fish was pooled as one sample, and six samples per treatment. The intestine samples were stored at -80°C to analyze the relative gene expression during a short-term exposure to the experimental feed.

2.3 Growth performance

At the beginning and end of the trial, fish in each tank were group weighed and counted to calculate growth performance, including Weight gain (WG %), Specific growth rate (SGR), Survival rate (SR %), PE (Protein efficiency %), Feed conversion ratio (FCR) (18). The production performance were calculated as follows:

Weight gain (WG, %) = $(FBW - IBW)/IBW \times 100;$

Specific growth rate (SGR, %/d)

$$= [ln(FBW) - ln(IBW)] \times 100 /d;$$

On the 56th day, after weight and length were measured, three fish from each tank (nine per treatment). The fish were then dissected to obtain the hepatopancreas and intestine quickly on ice. The hepatopancreas stored at -80°C for antioxidant capacity analysis. The intestinal tract was immediately divided into proximal intestine (PI), mid intestine (MI), and distal intestine (DI), stored at -80°C for key enzymes in tricarboxylic acid cycle and Na⁺/K⁺-ATPase analysis. An additional segment of PI, MI, and DI was fixed using Bouin solution for histology.

2.5 Antioxidant status

Glutathione (GSH, Cat. No. A006-2-1), malondialdehyde (MDA, Cat. No. A003-1) content; the activities of total superoxide dismutase (SOD, Cat. No. A001-1), and catalase (CAT, Cat. No.A007-1-1) were determined using the commercial kits purchased from Nanjing Jiancheng Bioengineering Institute.

2.6 Gene expression analysis

The total RNA was extracted from the proximal intestine, mid intestine, and distal intestine by Rapid Extraction Kit extracted total RNA (RN28; Aidlab Bio Inc., Beijing, China). RNA concentration and purity were measured by the spectrophotometry analysis (A260:A280 nm ratio) within the ratio specified by the kit (1.8-2.2). Subsequently, cDNA was synthesized by a MonScriptTM RTIII All-in-One Mix (MR05001M; Monad Bio Inc., Wuhang, China). The MonAmpTM SYBR[®] Green qPCR Mix (MQ10101S; Monad Bio Inc., Wuhang, China) used mRNA level analysis by quantitative real-time PCR on a CFX-96 Real-Time PCR Detection System (Bio-Rad Laboratories, Inc., USA). The reactions were done in a volume of 20 µL containing 0.4 µL of 10 µM each of reverse and forward primers, 1 μL of diluted cDNA, 10 μL of MonAmp TM SYBR $^{\textcircled{B}}$ Green qPCR Mix, and 8.2 µL of Nuclease-Free Water. PCR conditions were as follows: 95 °C for 5 min followed by 40 cycles of 95 °C for 10 s, 60°C for 10 s and 72 °C for 30 s. The online Primer3 (http:// primer3.ut.ee) software was used to design the primers. Target gene mRNA relative expression levels were calculated using the $2^{-\Delta\Delta Ct}$ method and corrected for the expression of the normalizing gene β actin, according to Luo et al. (19). The primers used for quantitative fluorescence analysis are shown in Table 2.

2.7 Histology of the intestine

The 1 cm segments were cut from the PI, MI, and DI. The 1 cm intestinal segments were processed, embedded, and stained according by Hangzhou Haoke Biotechnology Co., Ltd. Randomly selected fields were observed and pictures were taken by a camera coupled with a light microscope $(40\times)$. The images were measured and calculated for villus height, crypt depth, mucosal thickness according to the method previously described by Lin et al. (20) using the software Adobe Photoshop 2021. The average value was calculated from the measurement results from each sample index.

2.8 Key enzymes in tricarboxylic acid cycle and Na⁺/K⁺-ATPase in the intestine

Citrate synthase (CS, Cat. No. MM-204201), Isocitrate dehydrogenase (ICD, Cat. No. MM-3280701), α -ketoglutarate dehydrogenase complex (α -KGDHC, Cat. No. MM-9176301), and Na⁺/K⁺-ATPase (Cat. No. MM-166201), were determined by the enzyme-linked immunosorbent assays (ELISA), following the manufacturer's procedures (Jiangsu Meimian Industrial Co., Ltd., China)

2.9 Statistical data analysis

All statistical analyses were performed using SPSS, version 26.0 software. The level of significance was set to P< 0.05. Data were subjected to the one-way analysis of variance (ANOVA) followed by Tukey's test was performed for comparison of means, and data is presented as means ± standard error (SE).

3 Results

3.1 Growth performance

The effect of soybean antigen proteins and α -ketoglutarate on growth performance of mirror carp is shown in Figure 1. Compared with the FM group, there were no significant differences in SR. The WG, SGR, and PE were significantly reduced in SM and FMc group. In addition, the FCR was significantly increased in SM. Feeding the fish with the FMG diets did not affect significantly WG, SGR, SR, PE, and FCR.

Feeding the fish with the FMGA diets did not affect significantly WG, SGR, SR, PE, and FCR, when compared to the fish fed FMG diets. Feeding the fish with the FMcA diets did not affect significantly SR, PE, and FCR, but WG and SGR were significantly decreased, when compared to the fish fed FMc diets.

3.2 Antioxidant capacity in the hepatopancreas

The effect of soybean antigen proteins and α -ketoglutarate on antioxidative ability in the hepatopancreas was given in Figure 2. Compared with the FM group, there was no significant change in MDA content, however, SOD activity were significantly decreased in SM, FMG and FMc group; the CAT activity in FMG and FMc group and the GSH content in FMc group were decreased significantly.

Compared with the FMG group, there were no significant change in GSH content, MDA content, and CAT activity. However, the SOD activity of FMGA group increased significantly. Compared with the FMc group, there was no significant change in MDA content, CAT activity. However, the GSH content, SOD activity of FMcA group increased significantly. TABLE 2 Primer sequences of target genes from mirror carp (*Cyprinus carpio*), to measure the relative expression of the genes through quantitative real-time PCR.

Target genes	Primer sequence (5'-3')	Gene ID
TNF-α ^a	AAGTCTCAGAACAATCAGGAA TGCCTTGGAAGTGACATT	XM_019088899.2
TGFβ1 ^b	ACACGGTCACTTTGGTGTCA CAATGTGGGTTGCAGAACAG	XM_019072371.2
IL-1β ^c	AACTTCACACTTGAGGAT GACAGAACAATAACAACAAC	XM_019080073.2
claudin3c	GCACCAACTGTATCGAGGATG GGTTGTAGAAGTCCCGAATGG	XM_042710912.1
claudin7	CTTCTATAACCCCTTCACACCAG ACATGCCTCCACCCATTATG	XM_042732468.1
occludin	ATCGGTTCAGTACAATCAGG GACAATGAAGCCCATAACAA	KF975606.1
caspase3	CTCTACGGCACCAGGTTACTACTC GCCATCATTTCACAAAGGGACT	XM_019110173.2
caspase8	AAGCTCCTCATTGAAGAACCG ATCGTCCTGAACCACAACCTC	XM_042759183.1
caspase9	GCAAGCCCAAACTGTTCTTCAT CGTCCATCTGGTCATCTATCCC	XM_042750058.1
AMPKα ^d	GATGCCCTCTGGATGCTCTC GATGTCGTATGGTTTGCTCTGG	XM_042777568.1
АМРКβ ^е	AAACCTGAGGAACGCTTCAA CATCACATGGTTGGGTTCTG	XM_042711618
AMPKγ ^f	TGGGACAACAAACTGCAGAG GCAAGGGAATGAAGGAAACA	XM_042716989.1
ACC ^g	GTCACTGGCGTATGAGGATATT TCCACCTGTATGGTTCTTTGG	XM_042724983.1
TOR ^h	ATCTACGGCAAGACGAGAGG GTTGGTGGAGAGTGGGATCA	XM_042761449.1
4E-BP ⁱ	GGTGCTGATCAACGAGTCAA CAGGACGTTCTTGCTTGTCA	XM_042737175.1

^aTNF-α, tumor necrosis factor alfa;

^bTGF-β1, transforming growth factor beta;

CIL-1β, interleukin-1 beta;

^dAMPKα, AMP-activated protein kinase alfa;

^eAMPKβ, AMP-activated protein kinase beta;

fAMPKγ, AMP-activated protein kinase gamma;

^gACC, acetyl-CoA carboxylase;

^hTOR, target of rapamycin;

ⁱ4E-BP, eIF4E-binding protein.

3.3 Relative expression levels of inflammatory cytokines genes in the intestine

The effect of soybean antigen proteins and α -ketoglutarate on the mRNA levels of inflammatory cytokines genes in the intestines are shown in Figure 3. Compared with the FM group, the mRNA expression levels of TNF- α and IL-1 β were not significantly different, but the mRNA expression levels of TGF β 1 was significantly increased in SM and FMc group; the mRNA expression levels of TGF β 1, IL-1 β were not significantly different, but the mRNA expression levels of TNF- α was significantly increased in FMG group. Compared with the FMG and FMc group respectively, the mRNA level of IL-1 β was not significantly different in FMGA and FMcA group, but the mRNA expression levels of TGF β 1 were significantly increased in FMG and FMcA group, respectively.

3.4 Relative expression levels of junction protein in the intestine

The effect of soybean antigen proteins and α -ketoglutarate on the mRNA levels of junction protein in the intestines are shown in Figure 4. Compared with the FM group, the mRNA expression levels of claudin3c, claudin7, and occludin were not significantly different in SM and FMc group; the mRNA expression levels of claudin3c and occludin were not significantly different, but the mRNA expression levels of claudin7 was significantly decreased in FMG group.

Compared with the FMG group, the mRNA level of occludin was not significantly different, but the mRNA expression levels of claudin3c and claudin7 were significantly increased in FMGA group. Compared with the FMc group, the mRNA level of occludin and claudin7 were not significantly different, but the mRNA expression levels of claudin3c was significantly increased in FMcA group.

3.5 Relative expression levels of apoptosis factors in the intestine

The effect of soybean antigen proteins and α -ketoglutarate on the mRNA levels of apoptosis factors in the intestines was given in Figure 5. Compared with the FM group, the mRNA expression levels of caspase3, caspase8, and caspase9 were not significantly different in SM group. In FMG group, the mRNA expression levels of caspase3 and caspase8 were not significantly different, but the mRNA expression levels of caspase9 was significantly increased; the mRNA expression levels of caspase9 was not significantly different, but the mRNA expression levels of caspase3 and caspase8 were significantly increased in FMc group.

Compared with the FMG and FMc group, the mRNA expression levels of caspase3, caspase8, and caspase9 were not significantly different in FMGA and FMcA group.

3.6 Relative expression levels of AMPK/ ACC and TOR signaling pathway in the intestine

The effect of soybean antigen proteins and α -ketoglutarate on the mRNA levels of AMPK/ACC and TOR signaling pathway in the intestines was given in Figure 6. Compared with the FM group, the mRNA expression levels of AMPK α , target of rapamycin (TOR), and eIF4E-binding protein (4E-BP) were not significantly different, but the mRNA level of AMPK β , AMPK γ , and acetyl-CoA carboxylase (ACC) were significantly increased in SM group; the mRNA expression levels of AMPK β , ACC, TOR, and 4E-BP were



not significantly different, but the mRNA level of AMPK γ were significantly increased and the mRNA level of AMPK α were significantly decreased in FMG group; the mRNA expression levels of AMPK α , AMPK β , AMPK γ , TOR and 4E-BP were not significantly different, but the mRNA level of ACC was significantly increased in FMC group.

Compared with the FMG group, the mRNA expression levels of AMPK α , AMPK β , ACC, TOR, and 4E-BP were not significantly different, but the mRNA level of AMPK γ was significantly decreased in FMGA group. Compared with the FMc group, the mRNA expression levels of AMPK α , AMPK β , AMPK γ , ACC, TOR and 4E-BP were not significantly different.

3.7 Morphology of the intestine

The effect of soybean antigen proteins and α -ketoglutarate on morphology of the intestines was given in Figure 7. The fish fed on the SM diet, there were no significant change in crypt depth in DI; but the crypt depth were significantly increased in PI and MI. The villus height and mucosal thickness were significantly decreased in PI, MI, and DI. The fish fed on the SM diet, there were no significant change in villus height (MI) and crypt depth (DI); but the villus height (PI and DI) and mucosal thickness (PI, MI, and DI) were significantly decreased and crypt depth (PI and MI) were significantly increased. The fish fed on the FMc diet, the villus height and mucosal thickness were significantly decreased and crypt depth were significantly increased in PI, MI, and DI.

Compared with the FMG group, there were no significant change in villus height, crypt depth (MI and DI), mucosal thickness (PI and MI), but crypt depth (PI) was significantly decreased and villus height (PI), mucosal thickness (DI) was significantly increased in FMGA group. Compared with the FMc group, there were no significant change in villus height (PI, MI and DI), crypt depth (MI), mucosal thickness (PI and MI), but crypt depth (PI and DI) was significantly decreased and mucosal thickness (DI) was significantly increased.

3.8 Key enzymes in tricarboxylic acid cycle and Na^+/K^+ -ATPase in the intestine

The effect of soybean antigen proteins and α -ketoglutarate on key enzymes in tricarboxylic acid cycle and Na⁺/K⁺-ATPase in the intestines was given in Figure 8. Compared with the FM group, there were no significant change in CS, ICD, and α -KGDHC (PI and MI) activity, but Na⁺/K⁺-ATPase (PI) was significantly increased and α -KGDHC, ICD, CS (DI), and Na⁺/K⁺-ATPase



Mean values with different letters were significantly different (P<0.05).

(MI and DI) activity were significantly decreased in SM group; there were no significant change in CS, Na⁺/K⁺-ATPase (PI), ICD (MI), and α -KGDHC (PI and MI) activity, but CS, Na⁺/K⁺-ATPase (MI and DI), ICD (PI and DI), α -KGDHC (DI) were significantly decreased in FMG group; there were no significant change in ICD, α -KGDHC (PI and MI), and Na⁺/K⁺-ATPase (PI) activity, but CS (PI, MI and DI), α -KGDHC, ICD (DI), Na⁺/K⁺-ATPase (MI and DI) were significantly decreased in FMC group.

Compared with the FMG group, there were no significant change in CS, ICD, α -KGDHC and Na⁺/K⁺-ATPase (DI) activity, but CS, α -KGDHC, ICD and Na⁺/K⁺-ATPase (PI and MI)activity were significantly increased in FMGA group. Compared with the

FMc group, there were no significant change in CS (PI and MI), ICD, α -KGDHC (PI, MI, and DI), and Na⁺/K⁺-ATPase (PI and DI) activity, but Na⁺/K⁺-ATPase (MI) activity were significantly increased and CS (DI) activity was significantly decreased in FMcA group.

4 Discussion

In the present study, carp fed diets with soybean meal presented lower WG, SGR, and PE than in the control, which is corroborated by the findings of other studies investigating dietary soybean meal



Relative expression levels of inflammatory cytokins in the intestine of mirror carp at different treatments fed with diets containing fishmeal (FM), soybeans meal (SM), glycinin (FMG), B-conglycinin (FMC), AKG+B-conglycinin (FMCA) after eight weeks. TNF-a, tumor necrosis factor alfa; TGF-B1, transforming growth factor beta; IL-1B, interleukin-1 beta. Data are presented as means + SE, (n = 3). ^{a, b, c} Mean values with different letters were significantly different (P<0.05).



Relative expression levels of junction protein in the intestine of mirror carp at different treatment fed with the diets containing fishmeal (FM), soybean meal (SM), glycinin (FMG), B-conglycinin (FMcA)- AKG+B-conglycinin (FMGA), AKG+B-conglycinin (FMCA) after eight weeks. Data are presented as means + SE, (n = 3). ^{a, b, c} Mean values with different letters were significantly different (P<0.05).



FIGURE 5

Relative expression levels of apoptosis factors in the intestine of mirror carp at different treatments fed with diets containing fishmeal (FM), soybean meal (SM), glycinin (FMG), B-conglycinin (FMc), AKG+ glycinin (FMGA), AKG+B-conglycinin (FMcA) after eight weeks. Data are presented as means + SE, (n = 3). ^{a, b, c} Mean values with different letters were significantly different (P<0.05).



(21, 22). Dietary soybean meal increased the FCR, implying that the growth inhibition can be partly attributed to low feed efficiency by adding soybean meal (23). Glycinin and β -conglycinin are the main anti-nutritional factors in soybean meal, which can also adversely affect the growth performance of aquatic animals (3). In this study, dietary β -conglycinin reduced growth performance. This finding is consistent with studies in golden crucian carp (8). β -conglycinin cannot be digested and broken down into small molecules, and it is known to cause allergic reaction in intestinal tract, destruction of intestinal integrity, dysfunction of digestion and absorption of nutrients, inflammation, and oxidative damage, and finally suppressed fish growth performance (24). It was surprising to observe that growth performance was not influenced by the dietary Glycinin. These striking differences in growth performance between these two anti-nutrients may happen due to their functional properties. Glycinin has a higher content of sulfurcontaining amino acids than β -conglycinin (25). The composition of essential amino acids is more balanced, so it can better meet the body 's intake of essential amino acids. After adding the same dose of soybean antigen proteins, β -conglycinin could reduce the growth performance of carp (26), while glycinin did not affect the growth performance of turbot (27). Moreover, in this study fish were fed to apparent satiation (~9% of their body weight/day), and this high feed intake may have met the requirement of the growth performance of carps, which can be observed on the growth performance. Dietary AKG supplementation can promote growth performance by enhancing the activities of the antioxidant defense system (28). The supplementation of 1.0% AKG diet has also presented positive effects on growth performance of juvenile hybrid sturgeon (*Acipenser schrenckii* $\Im \times A$. *baerii* \Im). However, in this study, 1.0% AKG diet has not positive on the WG and SGR of carp, similar results were also observed in juvenile red drum (15). This lack of production performance may have been caused by the physiological differences among these fish species, breeding environments, growth phases, and dietary nutrient levels.

It is widely accepted that inflammation and oxidative stress usually occur simultaneously (29). Various stress factors may cause an imbalance between pro-oxidant and antioxidant systems, which will lead to oxidative stress. MDA, SOD and CAT were used as oxidative stress biomarkers (30). The concentration find in their tissues can reflect their antioxidant capacity (31) The oxidative stress induced by soybean antigen proteins, as evidenced by the depressed levels of SOD and CAT, indicates that glycinin and β conglycinin can cause SBM-induced oxidative stress in carp. These deleterious effects may be caused by the strong immunogenicity induced by soybean antigen proteins (32). Our results on the elevated activity of SOD suggest that AKG may suppress oxidative damage by glycinin and β -conglycinin. Besides these set of enzymes, GSH is one of the main non-enzymatic antioxidants that protect cells against oxidative damage (33), and it is protects many kinds of cells in body (34). AKG can be converted into glutamine by glutamate dehydrogenase and glutamine synthetase, which can be a marker of antioxidative function (14). It is evident that AKG can improve antioxidative capacity by increasing available glutamine and aiding antioxidative systems (14). Fish fed AKG supplemented feed presented an improvement of antioxidant enzyme levels, including GSH and SOD, indicating its protective efficacy (35). It has also been reported that AKG can positively alleviate the oxidative stress damage in cells, ultimately contributing to cell redox homeostasis (14), and also serve as an energy donor and antioxidant agent via the TCA cycle (36). The prooxidant effect of AKG seems to be due to the altered intensity of the TCA cycle in which AKG is preferentially metabolized (37). These data demonstrate that 1.0% AKG administration can alleviate the oxidative stress caused by the dietary soybean antigen proteins via the TCA cycle.

After feeding, soybean antigenic protein enters the intestine, destroying the structural stability of the intestine, activating the immune system and causing intestinal inflammation (38). TNF- α plays an essential role in mediating the inflammatory, regulating immune function and other pathophysiological reactions (39). IL-1 β is a pleiotropic inflammatory cytokine and plays an important role in the inflammatory process (40). Both TNF- α and IL-1 β are important pro-inflammatory factors (41). Intestinal inflammation is related to elevated pro-inflammatory cytokines (42). Enteritis is aggravated by up-regulating pro-inflammatory cytokines (43). In the present study, the expression of pro-inflammatory factor (TNF-



 α) was upregulated by dietary FMG group, which revealed the occurrence of glycinin-induced enteritis. Similarly, the phenomenon of glycinin-induced enteritis has also been observed in juvenile Rhynchocypris lagowskii (44). In addition, the mRNA level of TNF-a was decreased by dietary FMGA group. AKG can improve intestinal immunity and promotes intestinal health (45). Previously, Zhang et al. (34) found that glutamine suppressed TNF- α concentration. It is further confirmed by our study that AKG as a precursor for the biosynthesis of glutamine, increasing intestinal immunity. TGF- β is considered to be important anti-inflammatory cytokines that can inhibit inflammation (46). TGF- β has also played an important role in gut wound repair by promoting epithelial reconstitution through up-regulation of epidermal growth factor (47). Compared with the FMG/FMc group, the mRNA level of TGF-B1 was increased by dietary AKG. These data demonstrate that 1.0% AKG administration can alleviate the intestinal inflammation caused by the dietary soybean antigen proteins.

The intestinal epithelial barrier is the first line of defense against pathogenic microorganisms and toxins in the intestines (48). The connection between intestinal epithelial cells mainly relies on the TJ (49), and mainly composed of occludin, claudins, and others (50), which forms a highly selective barrier and protect animal health. In our study, we found that the mRNA levels of claudin7 was

suppressed by glycinin. Claudins are the major barrier-forming proteins of tight junction structure (51). Decreased gene expression of TJ usually represents the weakened intestinal barrier function (52). Similarly, Zhao et al. (53) reported that changes in TJ expression induced by soybean antigen protein lead to intestinal epithelial barrier dysfunction. It demonstrates that glycinin increase intestinal paracellular permeability, decrease tight junction proteins' expressions, suggesting that glycinin induce intestinal tight junction barrier dysfunction. The result in this study is supported by the findings in carp (54). This might be due to the fact that glycinin, as a foreign antigen, caused immune injury and inflammation, and then inflammatory factors destroyed intestinal epithelial tissue proteins (32). A lot of studies have proved that the impaired intestinal barrier is directly related to the intestinal inflammation (55, 56) and attenuate gut injury through an antiinflammatory role (57). In the study, the down-regulation of claudin3c, claudin7 observed after dietary soybean antigen protein was abolished by AKG. These observations were partly attributed to AKG supplementation improving glutamine synthesis activity and concentrations of glutamine and glutamic acid in the intestines (58). Dietary alanyl-glutamine in soy protein diets elevated the expression of TJ proteins (59). A number of studies have were reported that dietary glutamine enhance gut



development and regulate intestinal barrier function in multiple animal models (60). In this study, for the first time, demonstrate that 1.0% AKG administration can alleviate the damage of the epithelial barrier caused by the dietary soybean antigen proteins.

Apoptosis is a physiological programmed cell death through the activation of cell intrinsic suicide machinery (61). Apoptosis is a cascade amplifying reaction regulated by a cysteine family of proteases (62). It is well accepted that caspase family proteins is the central regulator of apoptosis (63). We found that the expressions of caspase3 and caspase8 were significantly elevated with FMc group. The expressions of caspase9 was significantly elevated with FMG group, consistent with the results of Peng et al. (64), who showed that soybean antigen protein increased the levels of caspase3 activity and induced piglet intestinal cell apoptosis. Additionally, glycinin or β-conglycinin accelerates enterocyte proliferation, apoptosis and migration for piglets (65). Hence this suggesting that soybean antigen protein causes intestinal damage. However, AKG administration could not alleviate the intestinal apoptosis caused by the dietary soybean antigen proteins in one week. Maybe the longer experimental period might have showed some positive on intestinal apoptosis.

AMPK, a sensor of the availability of intracellular energy, is activated at low energy levels and regulates cellular processes accordingly (66). AMPK is a heterotrimeric Ser/Thr kinase composed of an α , β and γ subunit (67). AMPK is activated by phosphorylation of Thr172 within the activation segment of the KD of the α subunit (68). The β subunits contains a region termed the carbohydrate binding module (68) and allows AMPK to interact with glycogen particles (69). AMPK preserves the level of ATP *via* stimulation of energy producing pathways and suppression of energy-consuming metabolisms (70). Under lowered intracellular

ATP levels, AMP or ADP can directly bind to the γ regulatory subunits of AMPK, leading to a conformational change that promotes facilitates the phosphorylation of AMPKa and the activation of AMPK (71). In the current study, the mRNA expression of AMPKy was significantly elevated and AMPKa was decreased with glycinin. This might be attributed to the fact that soybean antigen protein destroyed the structural stability of the intestine through pro-inflammatory and TJ proteins (54), which required more energy for maintenance, thus increased AMPKy gene expression. Maintaining the intestinal barrier is energy-consuming, and impairing mitochondria function is associated with intestinal inflammation (72). In addition, AMPK phosphorylation are reported to be involved in promote apoptosis (73). A similar phenomenon was also observed in the study that the expressions of TNF-a, caspase8, and AMPKy were significantly elevated and claudin7 was significantly decreased in FMG. It indicates that glycinin-induced injure intestine was partially related to the activate of AMPK signaling. In addition, the mRNA level of AMPKy was decreased in FMGA group. AKG as a source of energy (14), the oxidation of AKG through the tricarboxylic acid cycle can provide large amounts of ATP. AKG stimulates AMPK phosphorylation and oxidation of energy substrates in the intestinal mucosa, thereby enhancing ATP supply and supporting cell function (20). This suggested that AKG increased the intestinal ATP supply partly by down-regulating AMPKy mRNA expression. It indicates that AKG increases intestinal ATP supply by downregulating AMPKy gene expression, but the change of AMP/ATP is not enough to affect AMPK phosphorylation.

AMPK stimulates fatty acid oxidation as a way to increase energy levels. ACC is one of the first proteins identified as a target of AMPK (74). The mRNA level of ACC was increased with the soybean meal and β -conglycinin in this study. ACC is a critical enzyme of the lipogenic pathway (75); simultaneously, it is regulated by AMPK (76). This may indicate that dietary soybean meal and β -conglycinin could increase lipogenesis. In addition, the mTOR, it is also one of the target proteins of the downstream signaling pathway after AMPK (77). It is a limiting step in animal protein synthesis (78). And 4E-BP is one of the mTOR-regulated target sites. In this study, the mRNA level of TOR and 4E-BP were not significantly different with the SM, FMG and FMc for one week. However, studies have showed that β -conglycinin and/or glycinin could decrease the P-TOR protein expression for seven weeks in grass carp (24, 42). The mainly reason is unknown presently, maybe the longer experimental period might have showed some effect of the mTOR signaling pathway.

The fish growth highly depends on the healthy development of the intestine, which can be reflected by the villus height and mucosal thickness (9). Soybean allergy usually induces intestinal inflammatory diseases, characterized by the atrophy and proliferation of crypt villi, which accelerate the apoptosis and migration of intestinal cells (29). In this study, the VH and MT of the proximal intestine, mid intestine, and the distal intestine (except for the FMG group in MI) were decreased and crypt depth of the PI, MI, and DI (except for the SM, FMG group in DI) were increased in SM, FMG and FMc, indicates varying degrees of damage and inflammation. Similar results were also observed in turbot (8) and grass carp (79). Stimulation of antigen-active macromolecules can prompt an immune response that causes atrophy of the intestinal villi, thus damaging the intestinal morphology (9). In addition, dietary AKG has been reported to improve mucosal morphology and function of the intestine (57). In this study, the VH (PI) and MT (DI) were increased and crypt depth (PI) was decreased in FMGA group. The MT (DI) was increased and crypt depth (PI and DI) was decreased in FMcA, which was consistent with studies in piglets (80). In summary, soybean meal, glycinin and β-conglycinin damage the intestinal morphology and AKG improving intestinal morphology

Soybean antigen protein destroys the structural stability of the intestine and causes digestive and absorption disorders (54). Maintaining the intestinal barrier is energy-consuming, and impairing mitochondria function is associated with gut inflammation (72). Under stress, the body's energy metabolism disorders, ATP content decreased significantly, cell dysfunction or death, leading to damage to intestinal structure and function (81). Citrate synthase, isocitrate dehydrogenase and α -ketoglutarate dehydrogenase are the three rate-limiting enzymes of TCA cycle, which regulate the energy metabolism and biosynthesis of organisms. Citrate synthase can determine the rate of acetyl-CoA into TCA cycle, which is an important indicator of energy metabolism (82). Isocitrate dehydrogenase exists in mitochondria and cytoplasm and catalyzes isocitrate to produce α -ketoglutarate (83). The α -ketoglutarate dehydrogenase system exists in the mitochondrial matrix and catalyzes the oxidative decarboxylation of α -ketoglutarate to produce succinyl coenzyme A and reduced coenzyme I (NADH) (84). Na⁺/K⁺-ATPase mainly exists in the basement membrane of intestinal absorptive cells (85). It produces energy by decomposing ATP to maintain the reverse concentration

gradient of sodium and potassium. Its activity can indirectly reflect the absorptive capacity of intestinal mucosa (86). In this study, soybean meal, glycinin and β -conglycinin significantly decreased α -KGDHC, ICD, CS and Na⁺/K⁺-ATPase in DI, indicates that tricarboxylic acid cycle is inhibited and energy production is reduced. The AKG is an important intermediate product in the tricarboxylic acid cycle of organisms, and is also a biosynthetic precursor of glutamic acid, glutamine and arginine in organisms. In addition, glutamine and arginine play an important role in promoting the repair of damaged intestine (87). The AMPKy, VH (PI), MT (DI), α -KGDHC, ICD, CS, and Na⁺/K⁺-ATPase (PI and MI) were increased in FMGA. The Na⁺/K⁺-ATPase (MI) and MT (DI) were increased in FMcA. These data demonstrate that AKG in enterocytes may beneficially regulate the intracellular endogenous amino acid concentrations via TCA cycle (88) and then influence various signaling pathways (89), thereby alleviating the damage intestinal morphology caused by the dietary soybean antigen proteins. In addition, study has shown that AKG can be used as a substrate for intestinal energy metabolism (90). AKG could produce plenty of ATP in the TCA cycle and provide energy for intestinal cell processes (14). Dietary supplementation of AKG can improve the energy metabolism of intestine after soybean antigen protein stimulation.

5 Conclusions

In conclusion, dietary soybean meal can compromise the intestine health, and the adverse effects are related to the presence of β -conglycinin and glycinin, especially glycinin. 1.0% AKG may regulate intestinal energy *via* TCA cycle, thereby alleviating the damage intestinal morphology caused by the dietary soybean antigen proteins. Moreover, it is recommended the dietary supplementation of AKG when high levels of soybean meal are used in carp aquafeeds.

Data availability statement

The original contributions presented in the study are included in the article/supplementary material. Further inquiries can be directed to the corresponding author.

Ethics statement

This study was conducted following the China's Guidance of the Care and Use of Laboratory Animals. The experimental protocol was approved by the Committee on the Ethics of Animal Experiments of Huzhou University.

Author contributions

QL: project administration, writing the original draft preparation, and formal analysis; ZQ and RQ: formal analysis and investigation; FY: writing the manuscript and critical revision; YD and XL: investigation; QX: experiment design, management and manuscript revision. All authors contributed to the article and approved the submitted version.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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