

## Article

# Lotic Environment Affects Morphological Characteristics and Energy Metabolism of Juvenile Grass Carp *Ctenopharyngodon idella*

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**Abstract:** This study investigated the effect of a lotic environment on morphological characteristics and energy metabolism in juvenile grass carp *Ctenopharyngodon idella*. The fish were stocked in the lotic environment and forced to swim for 12 h per day for 4 weeks at three water current velocities of 0.5, 2, and 4 body length  $s^{-1}$  ( $Bl s^{-1}$ ). The control fish were stocked in the lentic environment with water current velocities of 0  $Bl s^{-1}$ . The results showed that lotic environment significantly increased body weight, body length, and condition factor of grass carp. The first principal component (PC1) characterized by measured overall body size suggested that fish in a lotic environment had body stoutness and wider tail stalk. Standard metabolic rate (SMR), maximum metabolic rate (MMR), and aerobic swimming performance ( $U_{crit}$ ) were elevated with the increased water flow and positively correlated with PC1. The 4  $Bl s^{-1}$  group showed significantly decreased contents of serum glucose and muscular glycogen, and a significantly increased level of serum lactic acid. The mRNA expression levels of AMP-activated protein kinase-phosphorylate PPAR  $\gamma$  coactivator 1  $\alpha$ -nuclear respiratory factor 1 (AMPK-PGC1 $\alpha$ -NRF1) pathway-related genes were significantly upregulated in red muscle of grass carp in the lotic environment. Water flow environment at 4  $Bl s^{-1}$  significantly increased ratios of metabolic enzymes (lactate dehydrogenase/citrate synthase) and cytochrome c oxidase/citrate synthase) in the muscle. The relationship between morphological characteristics and metabolic capacity suggested that the body size of grass carp in a lotic environment was shaped to promote energy metabolism. The study identified the evidence of the mechanism and relationship of the trade-off between energy and morphology in grass carp.

**Keywords:** lotic environment; energy response; metabolic regulation; AMPK pathway; *Ctenopharyngodon idella*



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## 1. Introduction

As an important factor in the hydrological environment, water flow has complex impacts on fish [1]. Previous research has proved that moderate water velocities have a range of positive effects on fish, including the enhancement of growth rate and disease resistance, while high water velocities tend to damage the fish's health and growth performance [2–4]. Fish are extremely susceptible to water velocity and swimming against the current because of their habits. The changes in the water environment can significantly affect fish ecology [5].

Swimming performance is an important indicator of whether fish can adapt to the environment, and is closely related to fish reproduction, escape ability, predation, migration, and environmental preferences [6,7]. Swimming performance has been often

used as a measure of ecological adaptability in fish. Different fish morphologies affect swimming performance [8,9]. In addition to swimming performance, energy metabolism also plays a vital role in the adaptation of fish survival in the environment [10,11]. The physiological and morphological changes of fish after swimming exercise are important manifestations of their adaptability [12,13]. In fact, the shape of fish is inseparable from their surroundings [14]. Environmental factors affect adaptive trade-offs in fish physiology, energy, behavior, and morphology [15,16]. Fish blood parameters have been considered as an indication of metabolism and health. Glycogen is the main storage form of glucose and the most important energy substrate during fish exercise [17]. However, there is limited acknowledgment on the trade-off mechanism between morphology and energy metabolism of fish in a lotic environment.

The AMP-activated protein kinase (AMPK) is a complex heterotrimer that is highly conserved in vertebrates, consisting of an  $\alpha$  catalytic subunit and two  $\beta$  and  $\gamma$  regulatory subunits [18]. Previous research has established that sustained swimming significantly increased AMPK expression level, thus facilitating downstream target gene expression in skeletal muscle in rainbow trout (*Oncorhynchus mykiss*) and promoting mitochondrial biogenesis and oxidative phosphorylation [19,20]. AMPK activation by hypoxic and low temperature in skeletal muscle also has several metabolic pathways, such as lipid oxidation and glucose glycolysis [21–23]. In the AMPK signaling pathway, peroxisome proliferator-activated receptor- $\gamma$  coactivator (PGC-1 $\alpha$ ) regulates the expression and function of the respiratory chain through nuclear respiratory factor-1 (NRF-1). Therefore, NRF-1 is an important downstream target of this pathway.

Grass carp (*Ctenopharyngodon idella*) possesses an important position in the aquaculture industry around the world. Prior studies that have noted the benefits of water velocity in swimming performance and growth of fish, but the mechanism of the effect of running water on the alteration of morphological characteristics and energy metabolism remains unclear in fish. This study aimed to investigate the role of the AMPK signaling pathway in grass carp stocked in the lotic environment, and to explore the relationship between morphological characteristics and swimming training. An integrated analysis of the metabolic capacity related to morphological characteristics of grass carp in the lotic environment was conducted to explore the relationship between metabolic and morphological characteristics. This research will provide data support for the physiological energetic and metabolic requirements of the adaptive responses to swimming in fish.

## 2. Materials and Methods

### 2.1. Fish and Experimental Conditions

Juvenile grass carp were acquired from a local fish farm (Wuqi Fishery in Wuhan, Hubei, China). Experimental fish were kept in the recirculating aquaculture system tanks (height 78 cm, diameter 1 m) for 2-week acclimation. The grass carp were fed with Tongwei commercial feed (crude protein  $\geq 31.0\%$ , crude fat  $\geq 4.0\%$ , crude fiber  $\geq 4\%$ , total phosphorus  $\geq 1\%$ , calcium  $\geq 0.5\text{--}2.0\%$ , sodium chloride  $\geq 0.3\text{--}1.2\%$ , moisture  $\geq 12.5\%$ , lysine  $\geq 1.4\%$ ) three times a day until apparent satiety. Water temperature and dissolved oxygen were 27–30 °C and  $> 7$  mg/L, respectively.

### 2.2. Experimental Protocol

After 2-week acclimation, 180 healthy grass carp (weight  $16.83 \pm 3.96$  g, body length  $9.43 \pm 0.86$  cm) were raised in 12 tanks. They were randomly assigned into one control group and three swimming training groups with water velocities of 0.5, 2, and 4 Bl/s, respectively. The water velocity of the control group was 0 Bl/s. The triplicates were used for each group. During the experimental period, the grass carp were forced to swim for 12 h (9:00–13:00, 14:00–22:00) per day in the custom-designed tanks. Each custom-designed tank was equipped with a cylindrical tube in the central area and a submersible pump at the bottom of the tank, as previously described by Zhang [24].

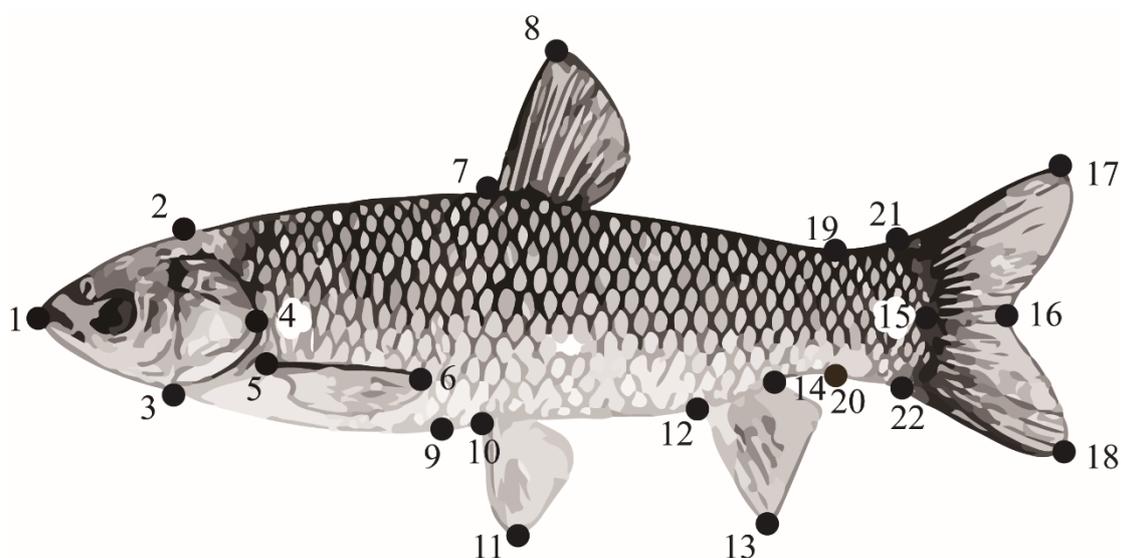
### 2.3. Sampling

At the end of the experiment, the final body weight and length of all fish from the tanks were measured to the nearest 0.1 g or 0.1 cm. Four fish of each tank were randomly selected to measure their morphometrics, metabolic rates, and aerobic swimming ability. Then, nine fish were randomly chosen per tank and anesthetized with 200 mg/L MS-222. Blood from the tail artery of each fish were clotted at room temperature for 30 min and then centrifuged at 3000 r/min for 30 min to get serum. Red and white muscles were isolated along the lateral line of each fish and snap-frozen in liquid nitrogen, then stored at  $-80^{\circ}\text{C}$  for mRNA and protein expression analysis.

### 2.4. Analytical Procedures

#### 2.4.1. Measurement of Morphology

Positions of the 22 points used for measurement of morphology are shown in Figure 1. Twenty morphological parameters (1–15, body length; 1–16, fork length; 1–17/18; total length; 1–2/3/4, head length; 2–3, head height; 2–7, back length; 7–9, body height; 7–8, dorsal fin length; 5–6, pectoral fin length; 10–11, pelvic fin length; 12–13, anal fin length; 17–18, caudal fin height; 17–21/18–22, caudal fin length; 19–20, caudal peduncle height; 14–22/19–21, caudal peduncle length) were measured by ImageJ. In order to eliminate the influence of individual size differences of grass carp on morphometric parameters, the standardized traits were obtained by dividing each morphological index by body length. On sampling day, the condition factor (CF) for each fish was calculated using the following formula:  $K = 100 \cdot \text{body weight} / (\text{body length})^3$ .



**Figure 1.** Data collection site diagram of physical characteristics in grass carp. Twenty-two points were set as the overall shape indicator in juvenile grass carp, as previously described in Lu [25].

#### 2.4.2. Measurement of Aerobic Swimming Ability

The grass carp were transferred into a custom-designed flow-through respirometer [26] and allowed to adapt for 1 h. During the test, the water velocity was then increased in  $1 \text{ Bl s}^{-1}$  increments from the initial value of  $1 \text{ Bl s}^{-1}$  every 20 min until the fish became fatigued. Water velocity and time were recorded.

$U_{\text{crit}}$  was calculated using Brett's equation:  $U_{\text{crit}} = V + (T/\Delta T)$ , where  $V$  is the maximum swimming speed ( $\text{Bl s}^{-1}$ ) of the fish in the whole cycle,  $\Delta T$  is the prescribed time (20 min) for each acceleration, and  $T$  is the time for the fish to swim to ultimate exhaustion (minutes).

#### 2.4.3. Measurements of Oxygen Consumption Rate

The oxygen consumption rate of the grass carp was also determined using a custom-designed flow-through respirometer. The oxygen content at the water outlets from each group was determined by a water quality analyzer (Hach 40H). At the end of the  $U_{crit}$  measurements, the fatigued grass carp was exposed to the air for 1 min [27] and then transferred into the flow-through respirometer to measure its maximum metabolic rate (MMR). The standard metabolic rate (SMR) of grass carp was quantified after 24 h. The metabolic rate was calculated using the following formula:

$$MO_2 = (\Delta O_2 \times V) / m^{0.75}$$

where  $\Delta O_2$  is the difference between the dissolved oxygen ( $mgO_2 \cdot L^{-1}$ ) in the breathing experimental chamber and the blank breathing chamber (without fish),  $V$  is the water velocity rate in the breathing chamber ( $L \cdot h^{-1}$ ),  $m$  is the weight of the fish (kg), and  $MO_2$  was adjusted to a standard body weight of 1 kg using a mass exponent of 0.75 [28]. The SMR was calculated by taking the average of the lowest three values of oxygen consumption rate during the 10-h measurement period.

#### 2.4.4. Determination of Contents of Lactic Acid and Glucose in Serum

Content of lactic acid in serum was measured with a colorimetric determination kit (JianCheng Bioengineering Institute, Nanjing, China). Glucose (GLU) content in serum was assayed by a Selectra-xl automatic biochemical analyzer (VITALAB, Dieren, Netherlands).

#### 2.4.5. Determination of Glycogen and Enzyme Activity of Muscle

Glycogen concentrations in muscle were detected by the sulfuric acid-anthrone colorimetric method according to the manufacturer's instructions (Nanjing Jiancheng) [29]. The lactate dehydrogenase (LDH) in muscle was quantified by LDH assay kit (Nanjing Jiancheng) [30]. Citrate synthase (CS) and cytochrome c oxidase (COX) Elisa kits were purchased from Jingmei, Shanghai, China.

#### 2.4.6. Gene Expression Level of AMPK Pathway in Red and White Muscle

The red and white muscle RNA were extracted with total RNA extraction reagent (Takara, Kyoto, Japan). Reverse transcription was performed using Hifair<sup>®</sup> III 1st Strand cDNA Synthesis Super Mix for qPCR (gDNA digester plus) (YEASEN, Shanghai, China). QPCR was performed with Hieff<sup>™</sup> qPCR SYBR<sup>®</sup> Green Master Mix (Low Rox Plus) (YEASEN) following the manufacturer's protocols in a Quant Studio<sup>®</sup> 6 Flex Real-Time PCR System (Life Technologies, Beijing, China) [31]. The relative expression of the target gene was determined by using the  $2^{-\Delta\Delta Ct}$  method for analysis and normalized using 18s RNA and  $\beta$ -actin. The gene-specific primer sequence reference is presented in Table 1.

**Table 1.** Primer sequences used for PCR analysis.

Gene	Forward (5'-3')	Reverse (5'-3')
AMPK $\alpha$ 1	TAGTGGCCGAGACTCAACCG	TCTGGCTACGGATTCCCAGG
AMPK $\alpha$ 2	AGTTCTACCTGGCCTCCAGT	GAGAGCATCCAGAGGGCAGC
AMPK $\beta$ 1a	TGGACAGCCCAGAAGATGC	TGGGTCGATCTAAAGTGAGGA
AMPK $\beta$ 1b	TGGAGCTCGACCCAAAATCC	AACACAGTGGGCCTTTCCTC
AMPK $\beta$ 2	GAGCCCGGCAAATGATGG	AGCCTGAGGACTCGTCTTT
AMPK $\gamma$ 1	AACACACTCTACATCCTGACAC	CCAGCGTTTTAGACAGAAAAGCA
AMPK $\gamma$ 2a	TTCAAGTCCAGCTGTTTGGT	GACGTCTCCAGCCAATAGCA
AMPK $\gamma$ 2b	CACGAGACCCTGGAGACCAT	TGCCATTCTGCATCTACCTG
AMPK $\gamma$ 3	CCCTCACGAAACCCTCGAAA	CAGAGAGCGAGACGATTCCC
PGC-1 $\alpha$	TCGGCCCCAGTCTCATTCTA	ATACACCACTCGCCTCTCCT
NRF-1	GTAGTGCAGACGATCAGTAATCC	GTCAGTGTGGAGTAATTCACCTGG
$\beta$ -actin	TGAAATTGCCGCACTGGTTG	CTGAGCCTCGTCACCAACAT
18sRNA	GGCGCGCAAATTACCCATT	TCCCGAGATCCAACACTACAAGC

### 2.5. Statistics

All analyses were carried out using SPSS 26.0 software (Chicago, IL, USA) and presented with mean  $\pm$  SEM. Normality tests were performed by using the Kolmogorov–Smirnov tests and all data were normally distributed. The morphology parameters were converted to percentage values of body length, and then data were converted with principal components analysis (PCA) and KMO and Bartlett’s sphericity test were performed, indicating that our data were adequate for factor analysis. Optimal factors were kept and rotated with an eigenvalue greater than 1, and used to analyze the relevant morphology, swimming performance and oxygen consumption rate by least squares linear regression. All data were analyzed by one-way ANOVA post-hoc Duncan test ( $p < 0.05$ ).

## 3. Results

### 3.1. Growth Performance

The fish growth was changed by the lotic environment for 4 weeks. The body weight, body length, and condition factor of the grass carp all increased significantly with the increase of water velocity at 0.5, 2, and 4  $\text{Bl s}^{-1}$  ( $p < 0.05$ ) (Table 2). The highest levels of body weight, body length, and condition factor occurred in the 4  $\text{Bl s}^{-1}$  group ( $p < 0.05$ ) (Table 2).

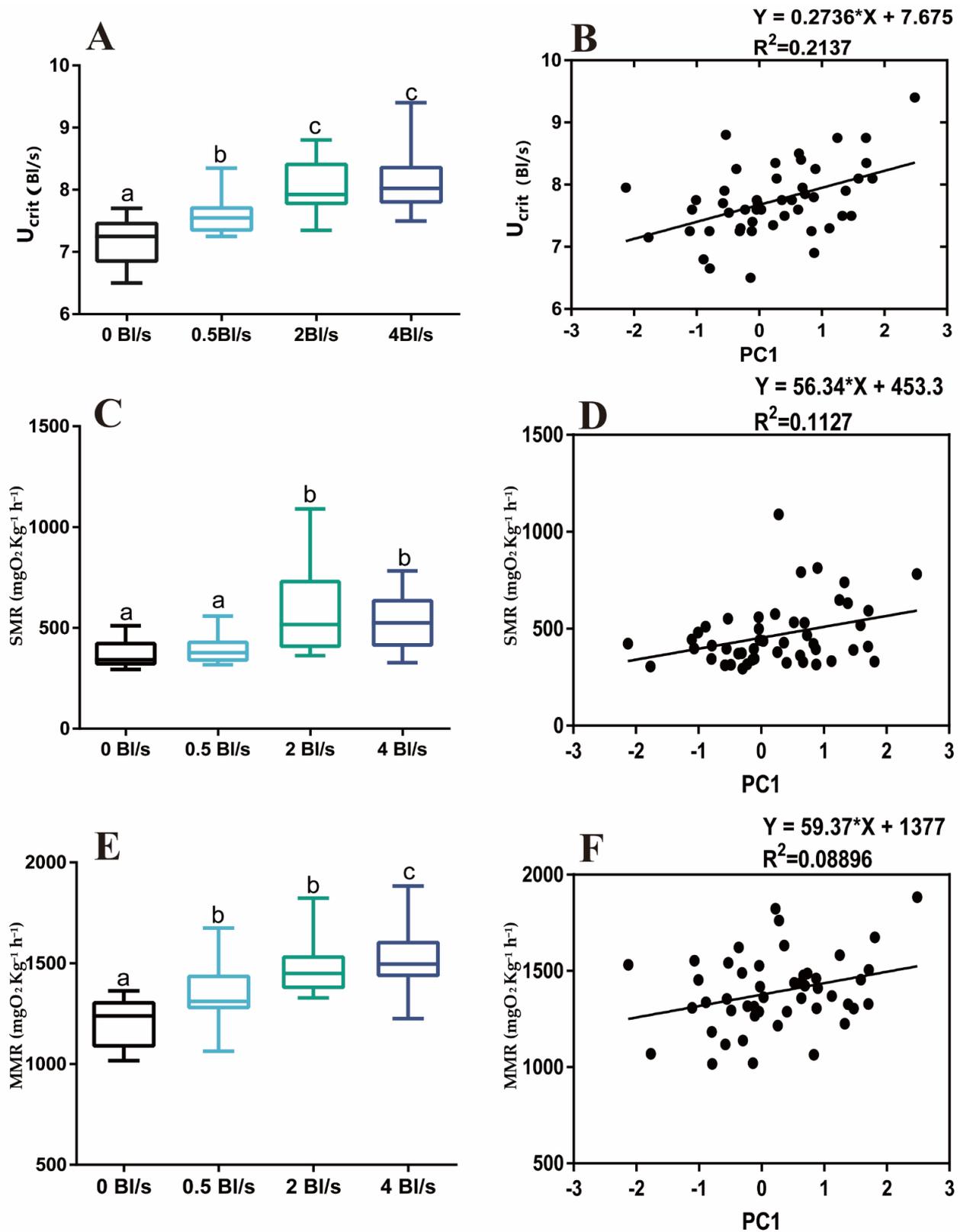
**Table 2.** Growth parameters of juvenile grass carp in the lotic environment.

Parameters	Water Velocity			
	0 $\text{Bl s}^{-1}$	0.5 $\text{Bl s}^{-1}$	2 $\text{Bl s}^{-1}$	4 $\text{Bl s}^{-1}$
Body weight (g)	23.08 $\pm$ 5.53 <sup>a</sup>	31.89 $\pm$ 13.78 <sup>b</sup>	31.72 $\pm$ 14.67 <sup>b</sup>	41.24 $\pm$ 18.91 <sup>c</sup>
Body length (cm)	10.38 $\pm$ 0.68 <sup>a</sup>	11.45 $\pm$ 1.38 <sup>b</sup>	11.78 $\pm$ 1.42 <sup>b</sup>	12.77 $\pm$ 1.36 <sup>c</sup>
Condition factor (CF)	1.95 $\pm$ 0.11 <sup>a</sup>	2.07 $\pm$ 0.18 <sup>b</sup>	2.08 $\pm$ 0.15 <sup>b</sup>	2.21 $\pm$ 0.21 <sup>c</sup>

Different superscripts indicated statistical significance for different training groups ( $p < 0.05$ ).

### 3.2. Morphological Characteristics, Aerobic Swimming Performance, and Oxygen Consumption Rate

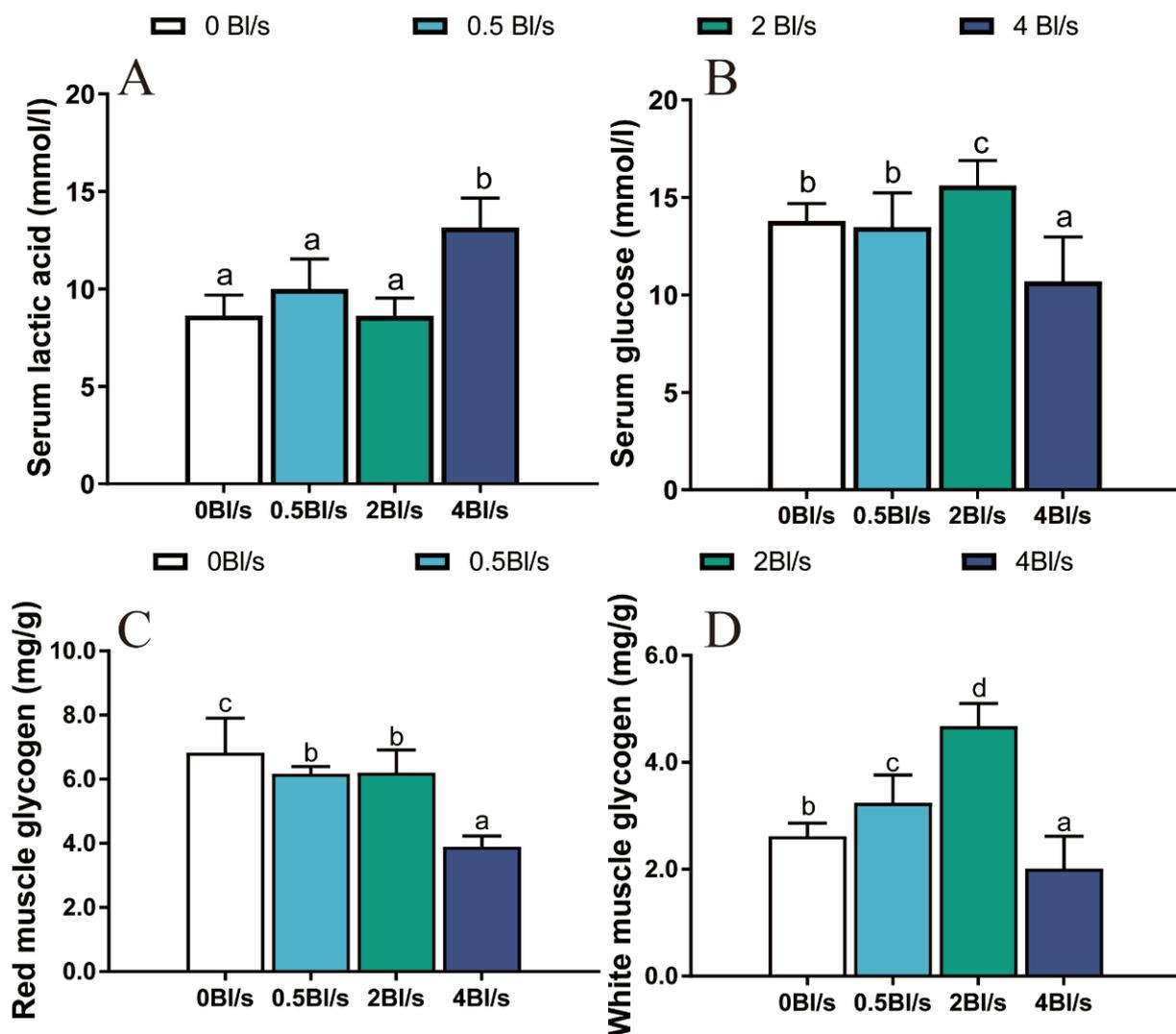
The water velocity significantly impacted the aerobic exercise capacity of grass carp. The  $U_{\text{crit}}$  of fish stocked in the lotic environment significantly increased compared with the control group ( $p < 0.05$ ) (Figure 2A). The diameters of the morphological characteristics fins, tail stalk and overall body stoutness were significantly increased after swimming (Table A1) ( $p < 0.05$ ). PCA showed that 19 morphological variables in this study produced six factors with eigenvalues greater than 1 describing 68.31% of the total variance. KMO and Bartlett of 19 morphological variables were KMO = 0.579  $>$  0.5, and  $p = 0.000 <$  0.05, indicating that the 19 morphological characteristics used in this study were suitable for principal component analysis. Subsequently, PCA was used with varimax normalized rotation to maximize the variances of the factor loadings across variables for each factor. Length of pectoral fin, head height, body height and caudal peduncle depth had high factor loadings for PC1. PC2 summarized the total length and caudal fin length and total length. PC3 showed head length. PC4 summarized the fork length and caudal fin height. PC5 was characterized by pelvic fin length and anal fin length. PC6 was characterized back length and dorsal fin length (Table A2). PC1 was positively correlated with  $U_{\text{crit}}$  ( $R^2 = 0.2137$ ,  $F_{1,46}$ ,  $p = 0.0009$ ) (Figure 2B). Among the six principal components,  $U_{\text{crit}}$  correlations were not significant for PC2, PC3, PC4, PC5, and PC6 (Table A3). The SMR of grass carp in the 2 and 4  $\text{Bl s}^{-1}$  groups were significantly higher than the control group ( $p < 0.05$ ) (Figure 2C). The MMR of 0.5, 2, and 4  $\text{Bl s}^{-1}$  groups were significantly higher than that of the control group ( $p < 0.05$ ) (Figure 2E). PC1 and metabolic rate were positively correlated (SMR,  $R^2 = 0.1127$ ,  $F_{1,46}$ ,  $p = 0.0197$ . MMR,  $R^2 = 0.08896$ ,  $F_{1,46}$ ,  $p = 0.0395$ ).



**Figure 2.** Linear regressions of PC1 by measurements of aerobic swimming ability and oxygen consumption metabolism in the lotic environment. (A) Aerobic swimming performance, (B) linear regressions of PC1 of aerobic swimming performance, (C) standard metabolic rate, (D) linear regressions of PC1 of standard metabolic rate, (E) maximum metabolic rate (F) linear regressions of PC1 of maximum metabolic rate. Different letters indicate significant differences between treatments ( $p < 0.05$ ).

### 3.3. Serum Biochemical Parameters and Muscle Glycogen Content

The content of LD did not significantly alter in either the 0.5 or 2 Bl s<sup>-1</sup> group compared with the control group, while it increased in the 4 Bl s<sup>-1</sup> group ( $p < 0.05$ ) (Figure 3A). Serum GLU content increased significantly in the 2 Bl/s group, but significantly decreased in the 4 Bl/s group compared with the control group ( $p < 0.05$ ) (Figure 3B). The glycogen content in red muscle was higher than that in white muscle in the same group. Compared with the control group, the contents of red muscle glycogen decreased significantly in the training groups ( $p < 0.05$ ) (Figure 3C). White muscle glycogen content increased significantly at 0.5 and 2 Bl s<sup>-1</sup> but was significantly decreased at 4 Bl s<sup>-1</sup> ( $p < 0.05$ ) (Figure 3D).

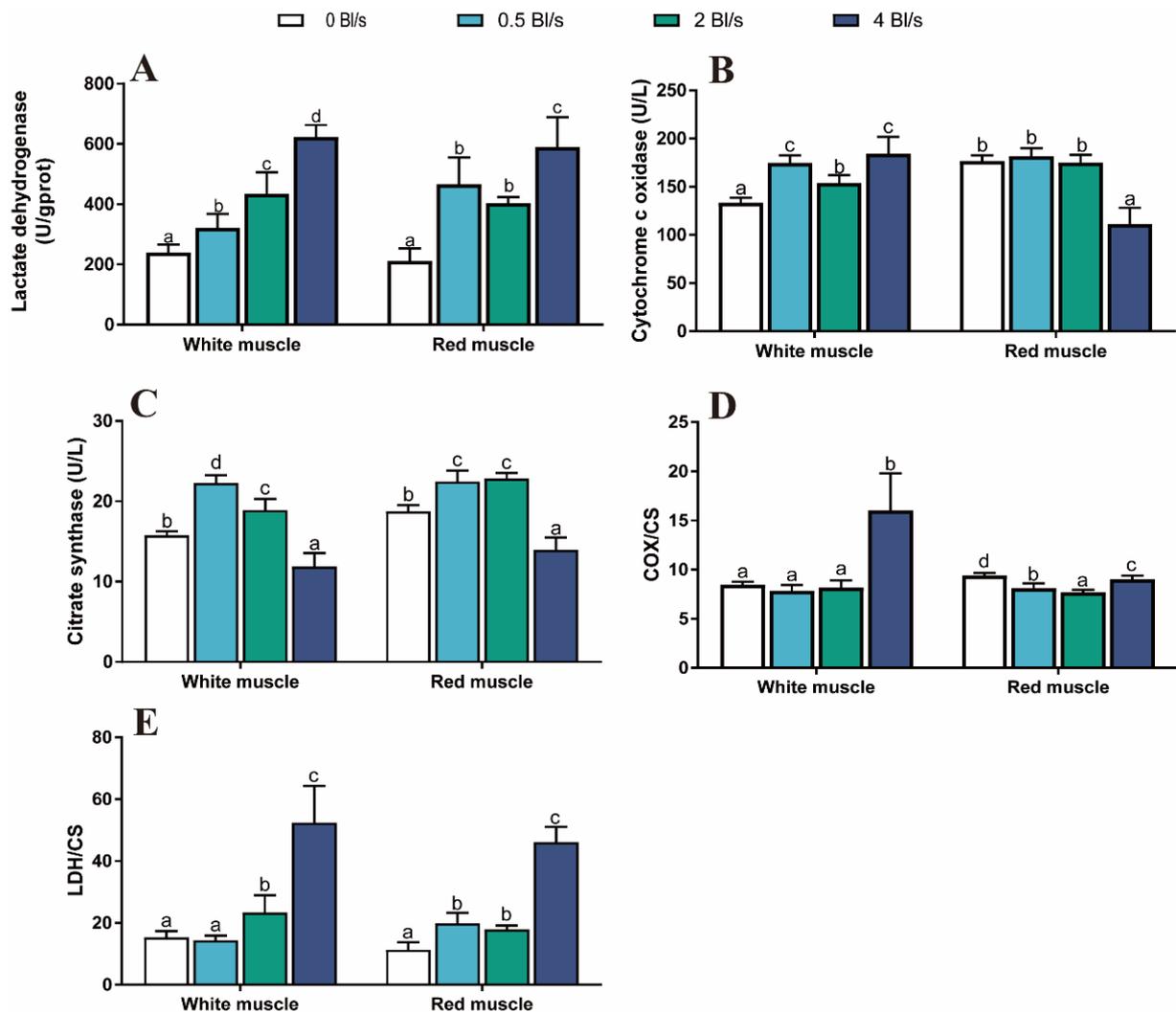


**Figure 3.** Effect of water velocity on lactic acid (A), glucose (B), glycogen of red muscle (C), glycogen of white muscle (D) in grass carp. Data are means  $\pm$  SEM ( $n = 9$  fish per treatment). Different letters indicate significant differences between treatments ( $p < 0.05$ ).

### 3.4. Muscle Enzyme Activity

The level of LDH activity was significantly increased in the lotic environment ( $p < 0.05$ ) (Figure 4A). The highest level of LDH was observed in the 4 Bl s<sup>-1</sup> group ( $p < 0.05$ ) (Figure 4A). The COX activity in white muscle increased significantly with the increasing water velocity ( $p < 0.05$ ) (Figure 4B). However, the COX activity in red muscle reduced significantly in the 4 Bl s<sup>-1</sup> group ( $p < 0.05$ ) (Figure 4B). Compared to the control group, the highest level of CS activity occurred in the 0.5 Bl s<sup>-1</sup> group, and the lowest level was found in the 4 Bl s<sup>-1</sup>

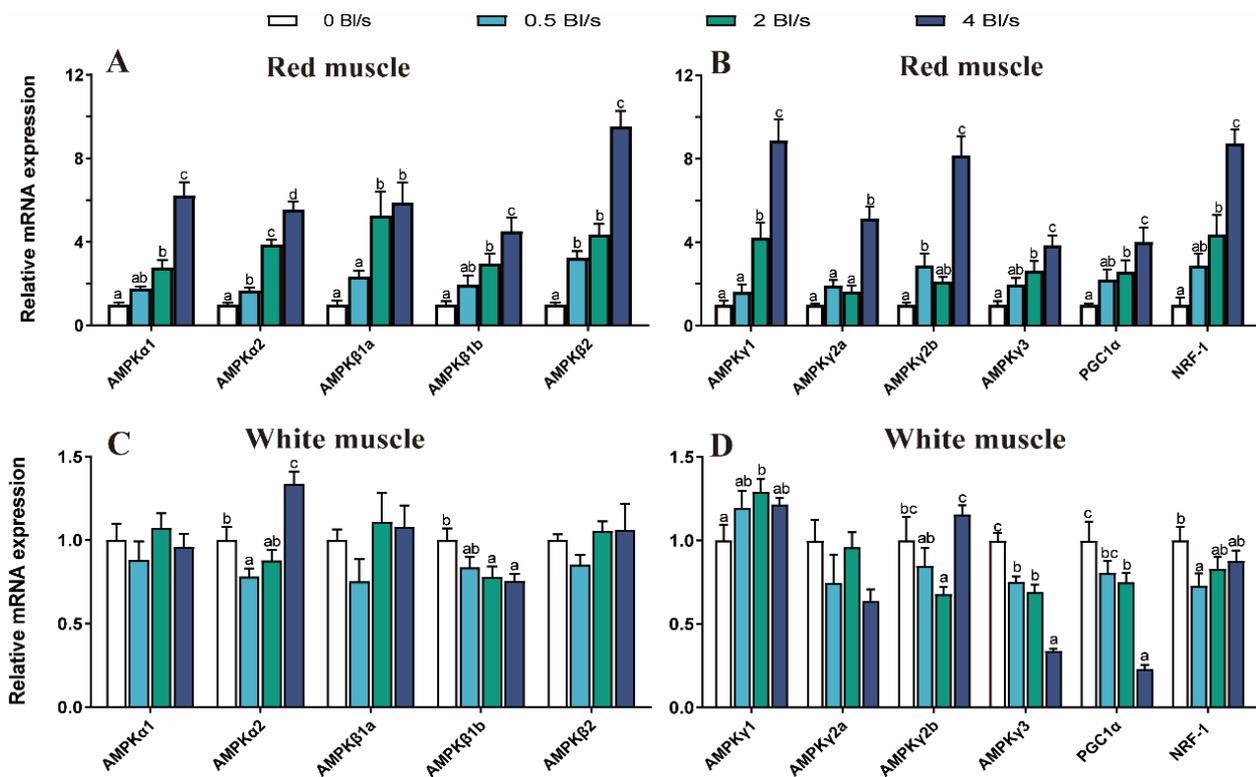
group ( $p < 0.05$ ) (Figure 4C). This trend was also found in red muscle (Figure 4C). The ratios of COX/CS and LDH/CS in the white muscle of the 4 Bl s<sup>-1</sup> group were significantly higher than those of the control group ( $p < 0.05$ ) (Figure 4D,E). In the red muscle, the COX/CS ratio significantly decreased in the 4 Bl s<sup>-1</sup> group ( $p < 0.05$ ) (Figure 4D).



**Figure 4.** Effect of water velocity on muscle enzyme activities of grass carp. (A) LDH activity in muscle. (B) COX activity in muscle. (C) CS activity in muscle. (D) COX/CS ratio in muscle. (E) LDH/CS ratio in muscle. Data are means  $\pm$  SEM ( $n = 9$  fish per treatment). Different letters indicate significant differences between treatments ( $p < 0.05$ ).

### 3.5. AMPK-PGC-1 $\alpha$ Signaling Pathway Gene Expression

The gene expressions of AMPK family, PGC-1 $\alpha$ , and NRF-1 were significantly upregulated in the exercise training groups compared with the control ( $p < 0.05$ ) (Figure 5A,B). In the white muscle, gene expression of AMPK $\alpha$ 2 was significantly upregulated in the 4 Bl s<sup>-1</sup> group compared with the control ( $p < 0.05$ ) (Figure 5C). In the swimming training groups, AMPK $\beta$ 1b, PGC-1 $\alpha$ , and NRF-1 mRNA levels significantly decreased with exercise training compared to the control group ( $p < 0.05$ ) (Figure 5C,D). The mRNA expression level of AMPK $\gamma$ 1 was increased in the lotic environment, and the expression in the 2 Bl s<sup>-1</sup> group was significantly upregulated ( $p < 0.05$ ) (Figure 5C). The AMPK $\gamma$ 2b mRNA expression level of the 2 Bl s<sup>-1</sup> group was significantly lower than the control group ( $p < 0.05$ ) (Figure 5C). There was no significant difference in AMPK $\alpha$ 1, AMPK $\beta$ 1a, AMPK $\beta$ 2, and AMPK $\gamma$ 2a mRNA expression levels among groups ( $p > 0.05$ ) (Figure 5C,D).



**Figure 5.** Effect of water velocity AMPK signaling pathway of grass carp in muscle. (A) AMPK $\alpha$  and  $\beta$  families mRNA expression levels in red muscle. (B) AMPK $\gamma$  family, PGC-1 $\alpha$  and NRF-1 mRNA expression levels in red muscle. (C) AMPK $\alpha$  and  $\beta$  families mRNA expression levels in white muscle. (D) AMPK $\gamma$  family, PGC-1 $\alpha$  and NRF-1 mRNA expression levels in white muscle. Data are means  $\pm$  SEM ( $n = 9$  fish per treatment). Different letters indicate significant differences between treatments ( $p < 0.05$ ).

#### 4. Discussion

The water current in the lotic environment forced grass carp to swim at the speed of the water velocity. Fish growth performance, metabolic rate, morphological and biochemical characteristics were changed by the lotic environment. Phenotypic plasticity of skeletal muscle is an indispensable part of the fish physiological response to swimming exercise [22]. In this study, the alterations in body weight, body length, and conditional factor appeared to be water velocity dependent. It is inferred that appropriate water velocity could stimulate the growth of grass carp. Body size is the main determinant of life history of animals, such as competitiveness, reproduction, and survival rate [32]. The principal component analysis showed that PC1 was mainly related to head height, fin length, and caudal peduncle depth, which were positively correlated with  $U_{crit}$  of grass carp. A previous study reported that deeper body shape increased hydrodynamic drag in fish during swimming [33,34]. This study found that the overall height was significantly increased, and the  $U_{crit}$  was also significantly increased in the lotic environment of grass carp. Other studies concluded that the deeper body shape of fish did not lead to decreases in locomotor performance [35]. When environmental conditions change, organisms are inclined to produce corresponding changes to maintain homeostasis [36]. Sustained water velocity seems to cause a trade-off in the morphology and metabolism of grass carp. “Trade-off” is understood as the balance between performance and adaptation when the organism is faced with two competing demands that cannot be simultaneously satisfied [37]. Exercise training reduced SMR, which reduced energy expenditure and expanded aerobic scope. By contrast, other studies have found that swimming training significantly increased the metabolic capacity of fish [38–40]. These studies showed that different species of fish have different energy metabolic varieties. In this study, water velocity significantly increased SMR and MMR of

grass carp, indicating that metabolic rate plasticity of grass carp includes an increase in oxygen consumption rate and metabolic capacity to adapt to the lotic environment [41,42]. An analysis of body shape and metabolism showed that PC1 had a significantly positive correlation with the metabolic capacity of grass carp. These findings suggested that water velocity significantly affected fish morphological characteristics and yielded a trade-off between body size and energy metabolism. Thus, the fish muscle would possess higher energy metabolism ability to overcome the increasing hydrodynamical drag.

The content of glucose and glycogen were significantly decreased in the 4 Bl s<sup>-1</sup> group, indicating consumption of carbohydrate reserves. In addition, the lactic acid content was also increased significantly in the 4 Bl s<sup>-1</sup> group, suggesting that 4 Bl s<sup>-1</sup> water speed increased the anaerobic exercise of grass carp. It is implied that higher intensity water speed caused more energy costs. This finding is in line with previous studies [43]. Glycogen is considered a storage form of energy and is related to swimming in fish [44]. The level of glycogen content in red muscle was higher than that in the white muscle of grass carp in the same water velocity training group. Consequently, red muscle provided much more energy in exercised grass carp. On this basis, we concluded that grass carp was involved in anaerobic metabolism at a water velocity of 4 Bl s<sup>-1</sup>.

AMPK has a direct role in regulating mitochondrial biosynthesis, muscle fiber formation, and endurance adaptability during long-term training [45,46]. In the present study, mRNA expression of AMPK-PGC-1 $\alpha$ -NRF-1 signaling pathway genes were significantly upregulated in red muscle. Furthermore, the expression of the AMPK family genes (AMPK $\alpha$ 1,  $\beta$ 1a,  $\beta$ 2,  $\gamma$ 2a) in white muscle were not significantly altered. The results showed that the responses of the AMPK pathway were fiber and training-intensity dependent, and in line with those of previous studies [47,48]. Moreover, lotic environment significantly upregulated AMPK $\alpha$ 2 mRNA in the 4 Bl s<sup>-1</sup> group. Fish AMPK $\alpha$ 2 isoform is involved in appetite regulation and is highly expressed in muscles [49,50]. As mentioned earlier, the 4 Bl s<sup>-1</sup> intensity water velocity increased the energy expenditure of grass carp. Previous studies found that exercise training produced higher appetite and food intake in fish [51]. PGC-1 $\alpha$  is the main one stimulating the energy metabolism in fish and plays an important role in various energy metabolism processes [22,52]. Exercise training improves mitochondrial biogenesis in muscles through PGC-1 $\alpha$ . In the present study, mRNA expression of PGC-1 $\alpha$  was significantly decreased in white muscle at 2 and 4 Bl s<sup>-1</sup> training groups, but significantly increased in red muscle at training groups. Studies suggested that the AMPK pathway was significantly increased at low temperature in Olive flounder (*Paralichthys olivaceous*), revealing its involvement in energy regulation [23]. Based on these findings, we speculated that the mRNAs of genes involved in mitochondrial functioning were more regulated in the red muscle than in the white muscle, and exercise training promoted energy metabolism in grass carp by increasing the AMPK pathway gene expression.

Mitochondria produce ATP through the process of aerobic respiration, involving the citric acid cycle, which can be adjusted in response to physiological stress [53,54]. In that respect, the CS enzyme involved in the citric acid cycle, is considered as a mitochondrial biogenesis marker. For energy metabolism, lactate dehydrogenase (LDH) and cytochrome c oxidase (COX) are important enzymes of anaerobic and aerobic metabolism, respectively. The levels of CS and COX were significantly increased after 0.5 and 2 Bl s<sup>-1</sup> water velocities, indicating that higher aerobic metabolism was induced in grass carp. This finding was also reported by McClelland [55]. The CS content was observed to be decreased in muscle of exercise-trained salmonids (1.5 Bl s<sup>-1</sup> for 4 weeks) [22], which indicated a complex and contradictory mechanism. Moreover, the COX activity was significantly increased in brown trout (*Salmo trutta*) treated with 1 Bl s<sup>-1</sup> water velocity but was significantly decreased in 2 Bl s<sup>-1</sup> water velocity exercised brown trout [56]. In the present study, high intensity water velocity (4 Bl s<sup>-1</sup>) significantly decreased the levels of CS and COX in red muscle, in agreement with the report by Blasco [57]. Moreover, in the present study, high intensity water velocity significantly increased the COX/CS ratio in white muscle. This indicated that the metabolic signatures of grass carp change under a lotic environment. In

that regard, the LDH/CS ratio was significantly increased in the 4 Bl s<sup>-1</sup> training group, which indicates that 4 Bl s<sup>-1</sup> water velocity treatment resulted in significantly enhanced anaerobic phenotype in grass carp to adapt to a high-intensity lotic environment. This observation may indicate that the mitochondria respiratory chain and tricarboxylic acid cycle (TCA) adapted the muscle metabolic flux towards to anaerobic metabolism from aerobic metabolism with the energy demand of the fish [58]. The findings of this study suggest that 4 Bl s<sup>-1</sup> intensity water velocity performed more anaerobic metabolism than that at 0, 0.5, 2 Bl s<sup>-1</sup> water speeds, and mitochondria of grass carp at 4 Bl s<sup>-1</sup> water velocity functionally adapted their energy consumption for swimming.

## 5. Conclusions

This study was designed to determine the mechanism of adaptability of a lotic environment on morphological and energy metabolism of grass carp. There was a trade-off between morphological characteristics and metabolic rate of grass carp in the lotic environment. The grass carp was involved in anaerobic metabolism at 4 Bl s<sup>-1</sup> and stimulated the transcriptional response of the AMPK pathway more in the red muscle. Water velocity of different intensities engendered different alterations in energy metabolism of grass carp. The juvenile grass carp could adapt to the lotic environment and enhance growth through energy metabolism and morphological changes. Based on the results, a water velocity of no more than 2 Bl s<sup>-1</sup> is recommended to apply to grass carp culture for better growth performance.

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**Data Availability Statement:** Data is contained within the article.

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

## Appendix A

### Appendix A.1. Differences in Morphological Characteristics

**Table A1.** Comparison of various indicators of grass carp in the lotic environment.

Character	Water Velocity			
	0 Bl/s	0.5 Bl/s	2 Bl/s	4 Bl/s
1–2/body length	23.82 ± 1.50	23.69 ± 1.11	23.38 ± 1.13	23.90 ± 1.16
1–3/body length	20.58 ± 0.97	19.95 ± 1.48	20.36 ± 1.21	19.90 ± 1.14
1–4/body length	25.47 ± 1.40	24.90 ± 1.09	25.07 ± 0.99	25.13 ± 1.12
1–16/body length	109.35 ± 1.09 <sup>b</sup>	108.53 ± 0.88 <sup>ab</sup>	108.46 ± 1.43 <sup>a</sup>	107.75 ± 1.57 <sup>a</sup>
1–17/body length	117.89 ± 1.59	117.49 ± 1.50	116.91 ± 1.76	117.15 ± 2.42
1–18/body length	120.01 ± 1.79 <sup>b</sup>	119.77 ± 1.54 <sup>b</sup>	118.91 ± 1.22 <sup>ab</sup>	117.90 ± 2.30 <sup>a</sup>
2–3/body length	18.94 ± 0.68 <sup>a</sup>	19.33 ± 1.02 <sup>a</sup>	19.10 ± 0.93 <sup>a</sup>	20.06 ± 0.90 <sup>b</sup>
2–7/body length	32.62 ± 2.49	32.60 ± 1.38	32.75 ± 1.85	32.31 ± 1.92

**Table A1.** *Cont.*

Character	Water Velocity			
	0 Bl/s	0.5 Bl/s	2 Bl/s	4 Bl/s
5–6/body length	15.32 ± 1.84 <sup>a</sup>	16.21 ± 1.08 <sup>ab</sup>	16.98 ± 1.49 <sup>bc</sup>	17.81 ± 1.00 <sup>c</sup>
7–8/body length	20.93 ± 1.94 <sup>ab</sup>	19.53 ± 2.24 <sup>a</sup>	20.62 ± 2.18 <sup>ab</sup>	21.54 ± 1.89 <sup>b</sup>
7–9/body length	23.22 ± 0.87 <sup>a</sup>	24.45 ± 1.52 <sup>b</sup>	24.72 ± 1.44 <sup>b</sup>	26.52 ± 2.16 <sup>c</sup>
10–11/body length	10.92 ± 1.15 <sup>a</sup>	11.09 ± 1.37 <sup>a</sup>	11.72 ± 1.29 <sup>ab</sup>	12.44 ± 1.28 <sup>b</sup>
12–13/body length	15.94 ± 2.40 <sup>b</sup>	14.21 ± 1.60 <sup>a</sup>	14.75 ± 1.73 <sup>ab</sup>	15.18 ± 1.65 <sup>ab</sup>
14–22/body length	12.48 ± 1.59	11.36 ± 2.01	12.03 ± 1.54	12.12 ± 1.48
17–18/body length	17.30 ± 4.22 <sup>a</sup>	20.48 ± 3.48 <sup>b</sup>	18.56 ± 5.36 <sup>ab</sup>	19.73 ± 3.81 <sup>ab</sup>
17–21/body length	21.49 ± 1.64	22.15 ± 1.37	22.07 ± 1.39	22.13 ± 1.98
18–22/body length	23.25 ± 1.42 <sup>ab</sup>	23.47 ± 1.23 <sup>b</sup>	22.93 ± 1.30 <sup>ab</sup>	22.48 ± 1.48 <sup>a</sup>
19–20/body length	10.25 ± 0.55 <sup>a</sup>	11.20 ± 0.81 <sup>b</sup>	11.00 ± 0.64 <sup>b</sup>	11.40 ± 0.96 <sup>b</sup>
19–21/body length	32.87 ± 2.40 <sup>a</sup>	33.56 ± 2.19 <sup>a</sup>	32.28 ± 2.65 <sup>a</sup>	35.34 ± 1.75 <sup>b</sup>

Different superscripts show statistical significance between groups ( $p < 0.05$ ).

*Appendix A.2. Principal Component Analysis*

**Table A2.** Principal component analysis of grass carp in the lotic environment.

Morphometric Characteristics	Principal Component 1	Principal Component 2	Principal Component 3	Principal Component 4	Principal Component 5	Principal Component 6
1–2/body length %	0.126	−0.066	0.835	−0.023	0.196	−0.077
1–3/body length %	−0.229	−0.077	0.714	0.011	−0.197	−0.042
1–4/body length %	−0.035	−0.130	0.904	0.101	0.067	0.093
1–16/body length %	−0.178	0.270	0.007	0.724	−0.091	−0.004
1–17/body length %	−0.055	0.220	0.035	0.755	0.024	−0.046
1–18/body length %	−0.179	0.774	0.017	0.367	−0.007	−0.144
2–3/body length %	0.616	−0.070	0.323	−0.364	0.022	0.045
2–7/body length %	0.035	−0.032	−0.088	0.424	−0.003	−0.778
5–6/body length %	0.832	−0.026	−0.051	0.177	−0.043	0.018
7–8/body length %	0.108	0.075	−0.091	0.279	0.182	0.757
7–9/body length %	0.843	−0.119	−0.054	−0.214	0.095	−0.007
10–11/body length %	0.334	0.002	−0.057	0.236	0.587	0.070
12–13/body length %	−0.425	0.244	0.111	−0.147	0.580	−0.117
14–22/body length %	−0.323	−0.013	−0.068	0.026	0.543	0.310
17–18/body length %	0.252	0.456	−0.247	−0.508	0.028	−0.257
17–21/body length %	0.014	0.821	−0.146	0.058	0.153	0.104
18–22/body length %	−0.065	0.851	−0.123	0.162	−0.155	0.139
19–20/body length %	0.747	0.058	−0.196	−0.334	0.088	−0.006
19–21/body length %	0.164	−0.078	0.093	−0.151	0.651	0.040
Characteristic value	2.981	2.449	2.318	2.208	1.591	1.433
Contribution rate	15.687545	12.88992	12.19804	11.61951	8.373102	7.541147
Cumulative Contribution rate	15.687545	28.57747	40.7755	52.39501	60.76812	68.30926

*Appendix A.3. Aerobic Swimming Performance and Morphological Characteristics*

**Table A3.** Linear regressions of PC by measurements of aerobic swimming performance of grass carp in the lotic environment.

Parameter	Principal Component	R <sup>2</sup>	p	Df	F
U <sub>crit</sub>	PC1	0.2137	0.0009	1,000, 46.00	12.5
	PC2	0.002314	0.7454	1,000, 46.00	0.1067
	PC3	0.00704	0.5707	1,000, 46.00	0.3261
	PC4	0.05543	0.1072	1,000, 46.00	2.7
	PC5	0.06287	0.0856	1,000, 46.00	3.086
	PC6	0.02617	0.2720	1,000, 46.0	1.236

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