



Mass scaling of the resting and maximum metabolic rates of the black carp

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Received: 2 November 2017 / Revised: 5 March 2018 / Accepted: 8 March 2018 / Published online: 22 March 2018
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Abstract

We investigated the body mass (M) scaling of resting metabolic rate (RMR), maximum metabolic rate (MMR), excess post-exercise oxygen consumption (EPOC), blood parameters, and organ masses of black carp (*Mylopharyngodon piceus*). The results showed that RMR scaled with M of the fish by an exponent (b) of 0.833 (b_R), which was significantly larger than 0.75. MMR scaled with M by a power of 0.775 (b_M), which was significantly lower than 1 and may be due to a small size proportion of red muscle. No difference between b_R and b_M or correlation between factorial aerobic scope and M was found. However, EPOC scaled positively with M by a power of 1.231, suggesting a constant aerobic capacity and an enhanced anaerobic capacity with fish growth. Mass of the inactive organs scaled with M by a power of 1.005, which was significantly larger than 1 and was negatively correlated with RMR, suggesting that the proportion of inactive organs increases with fish growth, which may contribute to the negative scaling of RMR. Red blood cell surface area (S) did not increase with increasing M , suggesting that the ontogenetic decrease in the surface area to volume ratio of cells may not contribute to the negative scaling of RMR. The predicted b_R value (0.846) by the average S (1.746 μm^2) differs by only 1.62% from the observed b_R value using our previously reported $S - b_R$ function in carp, suggesting that the species-specific cell size, rather than its ontogenetic change, affects the metabolic scaling of a species.

Keyword Oxygen consumption · Allometry · Body size · Metabolic level · Surface area

Abbreviations

M	Body mass
MR	Metabolic rate
RMR	Resting metabolic rate
MMR	Maximum metabolic rate

FAS	Factorial aerobic scope
EPOC	Excess post-exercise oxygen consumption
b	Scaling exponent
b_R	Scaling exponent of RMR
b_M	Scaling exponent of MMR
M_{O_2}	Oxygen consumption rate
Hb	Hemoglobin concentration
RBCC	Red blood cell count
S	Red blood cell surface area
$M_{\text{Active organs}}$	Total mass of active organs
$M_{\text{Inactive organs}}$	Total mass of inactive organs

Communicated by G. Heldmaier.

Electronic supplementary material The online version of this article (<https://doi.org/10.1007/s00360-018-1154-5>) contains supplementary material, which is available to authorized users.

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Introduction

The allometric scaling relationship between metabolic rate (MR) and body mass (M), a fundamental issue in eco-physiology, can be described by a power function $MR = aM^b$, where a is a constant and b is the scaling exponent. The mechanisms underlying metabolic scaling have attracted a large amount of research, and the allometric scaling of the metabolic rate has been suggested to be the consequence of

multiple causes (Glazier 2014a), such as the surface area-related heat loss (Rubner 1883), the ontogenetic decrease in the size proportions of active organs (Itazawa and Oikawa 1983; Oikawa et al. 1992), the fractal resource distribution network (West et al. 1997; Brown et al. 2004), the ontogenetic change in cell size and thus in the relative surface area of cells (Davison 1955; Kozłowski et al. 2003; Starostová et al. 2013), and the constraints resulting from the relative importance of metabolic level and effects of surface area vs. volume-related metabolic processes of organisms (the metabolic level boundaries hypothesis, MLBH) (Glazier 2005, 2010, 2014b). According to the aforementioned mechanisms, resting metabolic rate (RMR) can scale with M by various b values, e.g., 0.67 (Rubner 1883), 0.75 (West et al. 1997; Brown et al. 2004), and varying values between 0.67 and 1 (Davison 1955; Kozłowski et al. 2003; Glazier 2005, 2010).

Among most fish species, the b value for RMR (b_R) varies between 0.67 and 1 depending on lifestyle and environmental factors (Killen et al. 2007, 2010). Our recent studies quantifying metabolic scaling in several closely related species of cyprinids suggest that previous theories can only partially explain their metabolic scaling (Huang et al. 2013; Zhang et al. 2014; Luo et al. 2015). Data for additional species are necessary to test the previous theories of metabolic scaling. In recent work, we proposed a negative relationship between the red blood cell surface area (S) and b_R among cyprinid species. This can be explained by a smaller surface to volume ratio of the cells in species with larger cell sizes (Luo et al. 2015), which may intensify the surface boundary limits on b_R (Glazier 2005, 2010). It would be useful to validate the accuracy of this $S - b_R$ function in predicting the intra-specific b_R values by S in additional closely related species.

We selected the black carp (*Mylopharyngodon piceus*), a carnivorous cyprinid, as the focus of this study. Previous studies have reported the inter-individual variation in metabolic level, growth, and swimming performance of the black carp under multiple conditions (Yan et al. 2013; Pang et al. 2015, 2016). The metabolic scaling of this species and its underlying mechanism remain unknown. This study aims to assess the mass scaling of both RMR and MMR and the effects of cell size and organ size on metabolic scaling and to validate the $S - b_R$ function by Luo et al. (2015) in the black carp.

Materials and methods

Black carp specimens were collected from local fisheries in Xiema, Chongqing, China, and acclimated in a rearing system for at least 2 weeks prior to the experiment. During acclimation, the water temperature was maintained at

25 ± 1 °C with a 12 h:12 h light: dark photoperiod. The oxygen concentration was above 90% saturation, and the ammonia concentration was less than 0.015 mg L⁻¹. The fish were fed with a commercial diet once daily. The sample size of fish was 30. Animal handling and experiments followed the ethical requirements and recommendations of the Animal Care of the Fisheries Science Institution of Southwest University, China. All data generated or analyzed during this study are included in the supplementary information files (Online Resource 1).

When the acclimation was completed, the fish were starved for 48 h and weighed to 0.01 g. The metabolic rates of individual fish were measured by a continuous flow respirometer (Wang et al. 2012). According to the body mass of the fish, respiratory chambers of different sizes were chosen (0.03, 0.13, 0.52, 0.86, and 1.20 L). The fish were transferred individually into chambers, and the metabolic rate was measured after 12 h. Fourteen individuals were measured at the same time. One chamber without fish was set as a control. The flow rate was adjusted to ensure that the difference in oxygen concentration between the control chamber and the fish chamber was within 0.5–1.0 mg L⁻¹. An oxygen meter (HQ30D, HACH Company, Loveland, CO, USA) was used to determine the oxygen concentration of the chamber outlets. The following formula was used to calculate the oxygen consumption rate (M_{O_2} , mg O₂ h⁻¹):

$$M_{O_2} = \Delta O_2 \times v,$$

where ΔO_2 is the difference in the oxygen concentration (mg O₂ L⁻¹) between the fish chamber and the control chamber and v (L h⁻¹) is the flow rate through the chamber. M_{O_2} was measured hourly for 6 h, and the lowest two values was averaged as the RMR of the fish. The maximum metabolic rate (MMR, mg O₂ h⁻¹) was determined by the chasing protocol as described in Wang et al. (2012). After the RMR measurement, the flow rate was adjusted to ensure that 95% of the water in the chamber be refreshed in 1 min. The fish were transferred individually into the circle device and chased to exhaustion using a hand net (this process lasted for approximately 10 min). The fish were then rapidly returned to the respiratory chamber, and M_{O_2} was measured at 1 min intervals for the first 10 min post-exercise and then at 20, 30, 40, 50, 60, 80, 100, and 120 min until the M_{O_2} declined to within 1.2-fold of RMR. The peak M_{O_2} post-exercise was used as MMR. Factorial aerobic scope (FAS) was calculated as the ratio of MMR to RMR. Excess post-exercise oxygen consumption (EPOC, mg O₂) was calculated as the magnitude of excess oxygen consumption above RMR during the recovery phase.

After measuring the MMR, the fish were put into the anesthetic solution (MS - 222, 0.15 g L⁻¹) for blood and organ sampling. Blood was collected via caudal artery puncture using a 1 mL syringe containing 0.04 g L⁻¹

anticoagulant (1 g sodium fluoride:3 g potassium oxalate) (Huang et al. 2013). The Hb: hemoglobin concentration was measured with a spectrophotometer (752, Modern Science Company, Shanghai, China) at a wavelength of 540 nm. The red blood cell count (RBCC, 10^9 mL^{-1}) was measured by a Neubauer hemocytometer after dilution to 1:200 in 0.65% NaCl. The blood smear was dried and stained with Wright's–Giemsa fluid (Gao et al. 2007; Zhang et al. 2014). Photos were then taken using a microscope (EV5680B, Aigo Company, China). The red blood cell length (LC, μm) and red blood cell width (WC, μm) were determined using software (Image-pro Plus). The red blood cell was assumed as ellipsoid in shape, and the red blood cell surface area (S , μm^2) was calculated as follows: $S = LC \times WC \times \pi/4$.

Following blood sampling, the total mass of active organs ($M_{\text{Active organs}}$) was determined, including the brain, gills, red muscle, heart, hepatopancreas, digestive tract, spleen, gonads, kidneys, skin, and scales. The total mass of inactive organs ($M_{\text{Inactive organs}}$), including white muscle, abdominal adipose, fins, and bone, was calculated as the difference between M and $M_{\text{Active organs}}$.

Statistical analysis

The experimental data were analyzed using Microsoft Excel 2003 (Microsoft Corporation, Redmond, WA, USA) and SPSS 20.0 (SPSS Inc., Chicago, IL, USA). The log₁₀ transformed data of RMR, MMR, EPOC, S , Hb, RBCC, and M were used for the ordinary least square regression. Estimates of the scaling exponents were described with 95% confidence intervals (CIs). The differences between scaling exponents were analyzed using a general linear model (GLM) with M as a covariate. A t test was used to compare the difference between b values and 0.75 or 1. The relationships

between RMR and MMR, RMR and $M_{\text{Inactive organs}}$, RMR and S were analyzed using residual values. p values < 0.05 were considered statistically significant. Data are presented as the means \pm standard error.

Results

M of the experimental fish ranged from 3.7 to 202.1 g ($n = 30$). After exhaustive exercise, the metabolic rate values rose rapidly to a maximum followed by recovery to pre-exercise levels, and the recovery time ranged between 8 and 120 min dependent on M (Fig. 1). Whole animal RMR increased 21.9-fold with increasing M , ranging from 0.91 to 19.94 $\text{mg O}_2 \text{ h}^{-1}$, while MMR increased 21.5-fold, ranging from 6.92 to 149.12 $\text{mg O}_2 \text{ h}^{-1}$.

RMR and MMR scaled with M of the fish by powers of 0.833 (b_R) and 0.775 (b_M), respectively (Fig. 2). The b_R value was significantly larger than 0.75 ($t = 2.936$, $p = 0.007$), while b_M was not significantly different from 0.75 ($t = 0.592$, $p = 0.559$). Both b_R ($t = 5.950$, $p < 0.0001$) and b_M ($t = 5.277$, $p < 0.0001$) were significantly less than 1. There was no significant difference between b_R and b_M ($F = 1.264$, $p = 0.266$). No significant correlation was found between FAS and M ($r^2 = 0.074$, $p = 0.147$) (Online Resource 2). MMR and RMR were positively correlated when controlling for M (Fig. 3). EPOC increased with M by a scaling exponent of 1.231 (Fig. 4).

The scaling exponent value of the $M_{\text{Inactive organs}}$ (1.005) was significantly larger than 1 ($t = 2.568$, $p = 0.016$), while the scaling exponent value of the $M_{\text{Active organs}}$ (0.953) was not significantly different from 1 ($t = 1.949$, $p = 0.061$) (Fig. 5). RMR was negatively correlated with the $M_{\text{Inactive organs}}$ when controlling for M ($r^2 = 0.190$, $p = 0.016$) (Fig. 6).

Fig. 1 Metabolic rates of the black carp pre-exercise and post-exercise. Open circle: less than 10 g; filled circle: 10–30 g; open triangle: 30–90 g; filled triangle: larger than 90 g

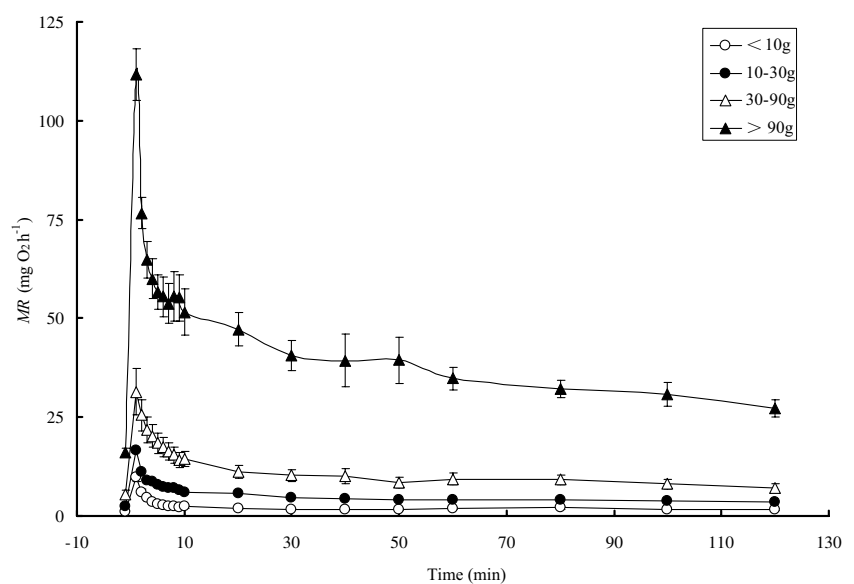


Fig. 2 Metabolic rate (MR) versus body mass (M) in the black carp. The data are logarithmic transformed to base 10. The open circles represent the maximum metabolic rate, MMR ($\text{mg O}_2 \text{ h}^{-1}$); the filled circles represent the resting metabolic rate, RMR ($\text{mg O}_2 \text{ h}^{-1}$)

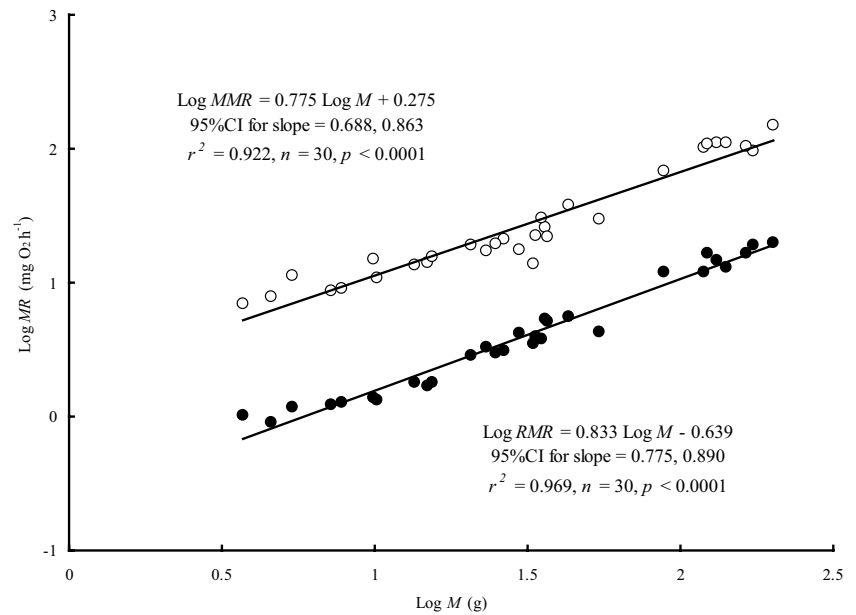
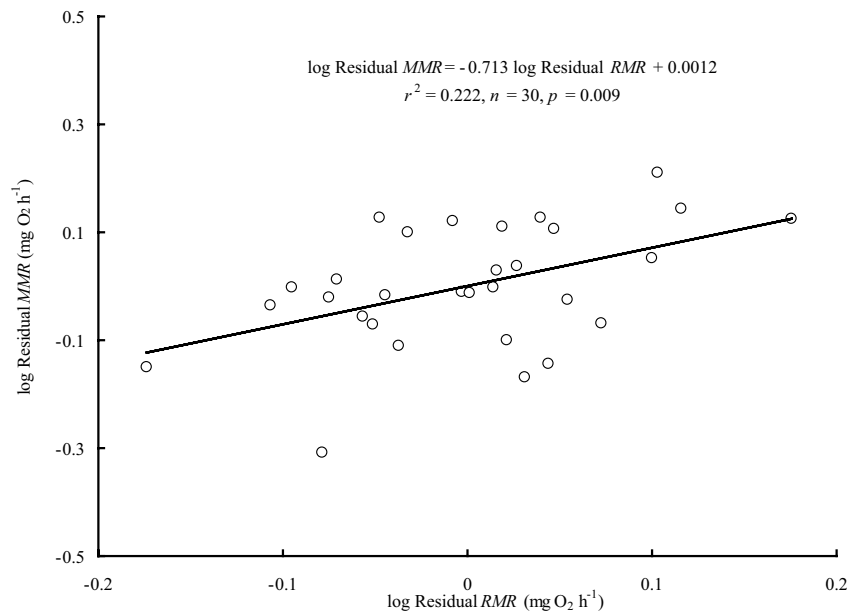


Fig. 3 Relationship between the residual resting metabolic rate (RMR, $\text{mg O}_2 \text{ h}^{-1}$) and the residual maximum metabolic rate (MMR, $\text{mg O}_2 \text{ h}^{-1}$) in the black carp



No obvious correlations were found between M and any of the blood parameters, including S ($r^2 = 0.0045, p = 0.726$), RBCC ($r^2 = 0.0454, p = 0.484$), and Hb ($r^2 = 0.0554, p = 0.221$) (Online Resource 3). No significant correlation between S and RMR was found when controlling for M (Online Resource 4).

Discussion

Our results indicate that the b_R value of the black carp is within the previously reported range of intra-specific scaling exponents (0.65–0.95) for most teleost fishes (Clarke and

Johnston 1999; Bokma 2004; Killen et al. 2010; Luo et al. 2015). Consistent with previous studies of several closely related carp species (Zhang et al. 2014; Luo et al. 2015), the b_R value of the black carp does not agree with the scaling exponent of 0.75 proposed by the fractal resource distribution network theory (West et al. 1997; Brown et al. 2004).

The cell size theory predicts that the b_R value should equal 1.0 if an increase in body size is entirely due to an increase in cell number rather than cell size and suggests that larger cells have relatively lower MR (Davison 1955; Kozłowski et al. 2003; Starostová et al. 2013). This theory has been supported by the negative correlations between RMR with S in the spined loach (*Cobitis taenia*) and the

Fig. 4 Relationship between the body mass (M , g) and the excess post-exercise oxygen consumption (EPOC, mg O_2) in the black carp. The data are logarithmic transformed to base 10

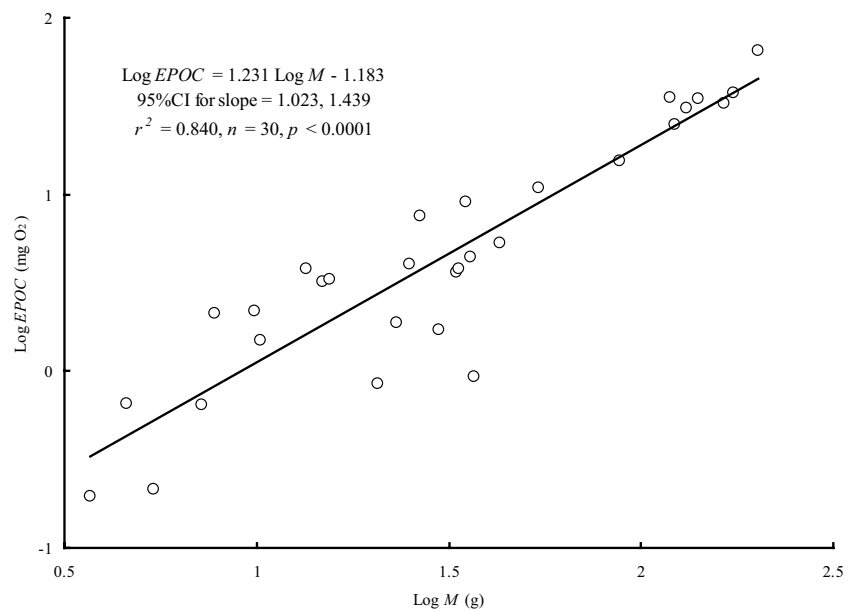
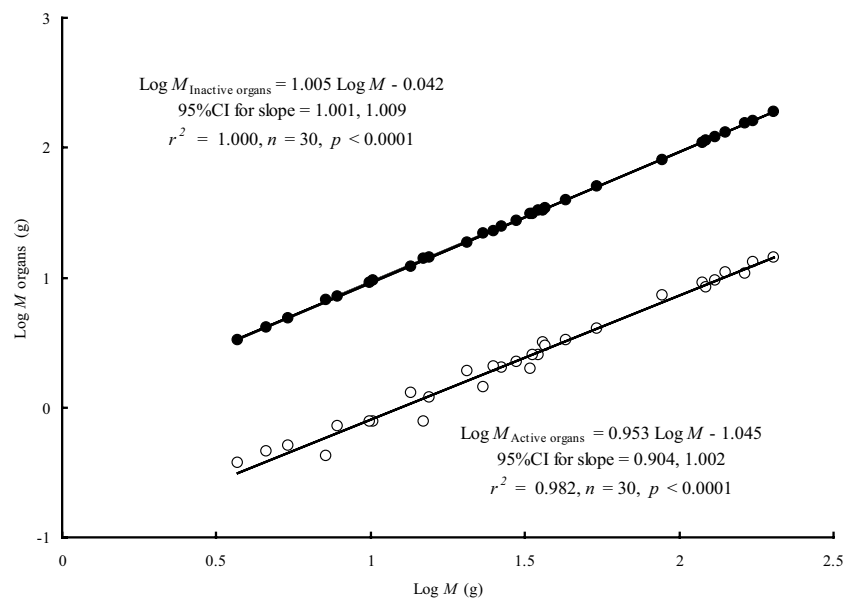


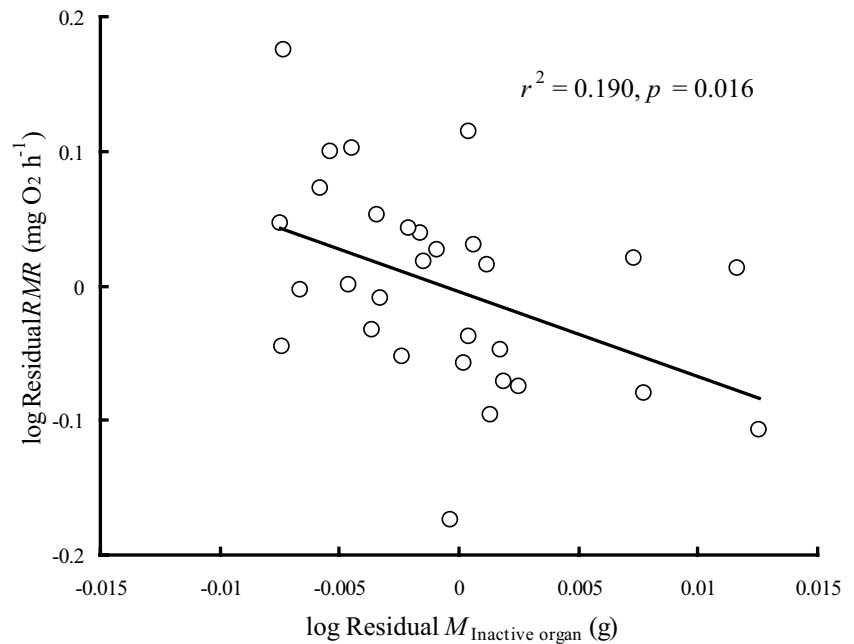
Fig. 5 Relationships between the masses of organs (M_{organs} , g) and the body mass (M , g). The data are logarithmic transformed to base 10. The open circles represent the active organ; the filled circles represent the inactive organ



crucian carp (*Carassius auratus*) (Maciak et al. 2011; Huang et al. 2013). However, no negative correlations between RMR with S were found in the black carp in the present study and several other species of carp in previous studies (Zhang et al. 2014; Luo et al. 2015), suggesting no general roles of cell size on RMR across different fishes. In addition, the present results showed that S of the black carp did not increase with increasing M (Online Resource 3), suggesting a constant cell size and thus a constant surface area to volume ratio of cells. Therefore, the ontogenetic decrease in the surface area to volume ratio of cells may not contribute to the negative scaling of RMR of the black carp. A similar unchanged erythrocyte size has also been reported in a

closely related species, the crucian carp (Huang et al. 2013). A potential explanation may be that MR is determined by factors other than ontogenetic change in the cell surface area to volume ratio, such as body size-dependent changes in the membrane permeability and metabolic activity of cells (Savage et al. 2007; Kozłowski et al. 2010). Alternatively, it has been proposed that the species-specific cell size, rather than its ontogenetic change, affects metabolic scaling. Species with a larger cell size may have a smaller surface area to volume ratio (Luo et al. 2015), which may intensify the surface boundary limits on b_R (Glazier 2005, 2010). Thus, a larger cell size results in a smaller b_R of a species. The negative relationship between b_R and S among several closely

Fig. 6 Relationship between the residual inactive organ mass ($M_{\text{Inactive organs}}$, g) and the residual resting metabolic rate (RMR, $\text{mg O}_2 \text{ h}^{-1}$) in the black carp



related species of carps can be expressed by $b_R = -0.260 \log S + 1.30$ (Luo et al. 2015). By this function, a b_R of 0.846 can be predicted by the logarithmic average S ($1.746 \mu\text{m}^2$) of the black carp, which differs by only 1.62% from the observed b_R value (0.833). The results suggest that the function described by Luo et al. (Luo et al. 2015) can effectively predict the intra-specific scaling exponent of RMR in carps.

According to the MLBH, a negative relationship exists between metabolic level (L) and b_R (Glazier 2005, 2010, 2014b). Killen et al. (2010) proposed an equation between the metabolic level and b_R among 89 species of teleost fish: $b_R = -0.145 \ln L + 1.377$. Using this equation, a recent study showed that the predicted b_R was only 7.5% deviated from the observed value in the crucian carp (Huang et al. 2013). In the present results, the natural logarithmic mass-specific RMR ($4.88 \text{ mg O}_2 \text{ kg}^{-1} \text{ h}^{-1}$) of the black carp at the mid-point of the regression ($\ln M = 3.31$) predicts a b_R of 0.669, which is 19.6% deviated from the observed b_R (0.833). The results cannot be explained according to the MLBH as that the black carp has an intermediate metabolic level in compared to that of those teleost species, which should entail an intermediate metabolic scaling exponent within the limited range (0.67–1) (Glazier 2005; 2010, 2014b).

It had been suggested that the allometric scaling of MR is attributed to a gradual decrease in the proportion of the $M_{\text{Active organs}}$ while increasing the proportion of the $M_{\text{Inactive organs}}$ with increasing M (Itazawa and Oikawa 1983; Oikawa et al. 1992). Consistently, the positive scaling of $M_{\text{Inactive organs}}$ with increasing M of the black carp suggests that the proportion of $M_{\text{Inactive organs}}$ increases as the fish grows (Fig. 5), which can contribute to the negative scaling of RMR of the black carp. In addition, the negative

correlation observed between $M_{\text{Inactive organs}}$ and RMR suggests that the variation of RMR among individuals may be due to intra-specific differences of $M_{\text{Inactive organs}}$ (Fig. 6). Because the scaling exponent of $M_{\text{Inactive organs}}$ (1.005) was very close to 1 and the r^2 for the correlation between RMR and $M_{\text{Inactive organs}}$ was small (0.19), this suggests that the intra-specific metabolic scaling of the fish can only partly be explained by the size changes of organs, and therefore other organ characteristics (e.g., organ metabolic level) may be involved.

The b_M value of the black carp was less than 1, suggesting that MMR may not scale isometrically with M . In many salmonids, MMR scales approximately isometrically (Brett 1965; Wieser 1985), which is attributed to the increasing importance of volume-related muscular energy expenditure on metabolism during exercise and the linear increase in muscle mass in proportion to M (Glazier 2005, 2009). However, in several species of carp, MMR scales negatively with M , suggesting that muscular energy expenditure has a limited contribution to whole-body metabolism in carp (Huang et al. 2013; Zhang et al. 2014; Luo et al. 2015). A possible explanation is that only a small portion of the carp's body contains red muscle. Indeed, the maximal locomotion speeds of carp are also very low (Yan et al. 2013).

The difference of b_M and b_R may determine a change in FAS, a species aerobic capacity-related parameter, as organisms grow. The lower b_M value than b_R value of the black carp results in no significant change in FAS with increasing M (Figs. 2, S1). This suggests that, in contrast to the general observations in salmonids (Brett 1965; Beamish 1978), the aerobic capacity of the black carp may not increase as their bodies grow. Accordingly, several oxygen capacity-related

blood parameters, including Hb, RBCC, and *S*, are also independent of *M* (Online Resource 3). In contrast, we found that EPOC scaled positively with *M* by a scaling exponent of 1.231 (Fig. 4), suggesting a higher metabolic consumption level restoring the RMR physiological state in larger fish.

MMR of the black carp correlated positively with RMR after controlling for body size (Fig. 3). This is consistent with the intra-specific results of the same species (Pang et al. 2015), as well as the inter-specific results among teleost fish species (Killen et al. 2016) and the intra- and inter-specific results of vertebrates (Auer et al. 2017), suggesting that there is a cost of maintaining the machinery that supports the maximal aerobic capacity. Similar positive relationships between RMR and MMR were found in the rainbow trout (*Salmo gairdneri*) (Wieser 1985), the brown trout (*Salmo trutta*) (Norin and Malte 2012), the crucian carp (Huang et al. 2013), and the grass carp (*Ctenopharyngodon idellus*) (Zhang et al. 2014).

In conclusion, the present study quantified the allometric scaling of the metabolic rate of black carp. The results suggest that the negative intra-specific scaling of RMR of the fish can be partly explained by the size changes of active/inactive organs. It suggests that mass-specific metabolic rate of active organs likely decreases with body mass.

Acknowledgements This work was supported by the National Natural Science Foundation of China (no. 31672287) and the Start-up Research Grant from Qinzhou University (no. 2017KYQD107). We thank Mr. Bo Zhang for his help in fish collection and thank the anonymous reviewers for their comments on the manuscript.

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