

Cooling of cultured water can resist heart failure caused by collagen deposition and necrosis of cardiac fibers under chronic heat stress in sturgeon

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Research

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Abstract

Background

Chronic heat stress (CHS) may threaten the survival of cultivated and wild sturgeon by jeopardizing heart function.

Methods

Therefore, we established a heart damage model for the Siberian sturgeon (*Acipenser baerii*) using varying degrees of CHS (24°C and 28°C) to explore its effect on the heart structure and function, and their mutual relationship.

Results

Our research showed that CHS caused systemic heart failure in *A. baerii*, clinically manifested as severely irregular ventricles, increased myocardial fibrotic in the interstitium of cardiomyocytes, and myocardial necrosis. Echocardiographic imaging of *A. baerii* revealed an accelerated heart rate, incomplete ventricular contraction, decreased cardiac output, and significantly reduced pumping efficiency under CHS. Generally, the contractility of the heart decreased and the afterload increased under CHS, which is typical of high-resistance and low-output heart failure. However, cooling of cultured water (20°C) can offset the adverse effects of partial or total CHS on tissue structure and function.

Conclusion

Our results systematically characterized the relationship between the effects of CHS on the heart structure and heart function in sturgeon. This work provides a preliminary reference for future summer breeding pond management and protection of sturgeon.

Introduction

Global warming and the construction of river dams can cause the water temperature of rivers to increase for extended periods of time which threatens the survival of aquatic organisms [1-6]. Chronic heat stress (CHS) occurs when the environmental temperature exceeds the optimal temperature range for fish to survive for a long-term [7-10]. CHS can directly threaten the physiological stability of freshwater species, affecting their behavior, growth, development, reproduction, immune, digestion, and circulatory system, and can even cause death [11-16].

Acipenser baerii is widely distributed in important economic fish in the northern hemisphere, and the caviar produced by adult fish has extremely high nutritional value [17–20]. Sichuan Province's sturgeon

farming production ranks among the top in China, and its normal growth temperature of sturgeon is maintained at 5-20°C all year round (Data form Sichuan Runzhao Fishery Co., Ltd). However, in summer, the water temperature of sturgeon breeding ponds often increases suddenly and lasts for 9-12 days, causing the CHS and a large number of deaths of cultured sturgeons. Mai et al. found that the growth of *A. baerii* was significantly hindered when the feeding water temperature was maintained at 24°C, and the survival rate at 27°C was significantly reduced [21]. Our team's CHS experiments show that the survival of *A.baerii* decreases when the water temperature exceeds 20°C (Yang et al. 2021). In the process of growth and development, the normal maintenance of heart function is essential to ensuring normal growth, development, and reproductive behavior [22]. Heart, the life-sustaining organ of fish, generates pressure to circulate the blood, transport oxygen from fish gills to various tissues, and remove metabolic waste products, thereby supporting active biological processes [23]. Therefore, the normal physiological temperature of the heart is the prerequisite for maintaining the high production efficiency of sturgeon. In recent years, an increasing number of studies have examined the effects of heat stress on sturgeon [6, 14, 24]. However, few have examined the effect of CHS on the sturgeon heart and establish the relationship between heart structure and function.

Tissue blood perfusion requires rhythmic and periodic contraction and relaxation of the heart ventricles and complete cardiomyocyte support. Among them, the changes in ECM (extracellular matrix), led by fibrillar collagen composition, play an important role in affecting the heart movement of fish[25]. Fibrillar collagen is the most common structural fibres in the ECM. The fibrillar collagen forms stiff fibres that maintain and support the alignment of myocardial cells by bearing wall stress[26, 27]. However, excessive deposition of cardiac collagen will increase the passive stiffness of the chamber wall and reduce the chamber compliance and dilatability of the ventricular cavity, which can have implications for diastolic filling[28]. However, changes in the composition of myocardial collagen in the heart of *A. baerii* under heat stress and the relationship between the composition of myocardial collagen and changes in cardiac function need to be further studied.

Moreover, pumping blood is considered the main function of the heart, its performance can affect fish behavior, as well as changes in organs, tissues, and cells (intercellular stroma) [15, 29–31]. However, adverse and extreme external pressure stress may cause damage to the structure of myocardial cells and cause necrosis [32]. This is particularly concerning because most fish hearts lack coronary circulation and only receive oxygen via diffusion from the venous return. Unlike the spongy layer, which receives venous blood, the compact layer of the fish ventricle has a coronary blood supply from the gill [33]. However, heat stress can cause serious damage to the gill tissue of fish, decrease in oxygen exchange capacity, and severely damage the oxygen supply to the heart [34]. For fish, the thickness and structural integrity of the dense layer of the ventricle are important prerequisites for maintaining the pumping function of the heart [35, 36]. This heart organ organization can impose limits on sustaining cardiac function at different temperatures and may lead to myocardial cell necrosis[37]. However, the influence of *A. baerii* myocardial necrosis under CHS on cardiac structure and the relationship between myocardial necrosis and changes in cardiac function need further research.

In present study, we systematically investigated the behavior of sturgeon under CHS, as well as the structure, function and molecular regulatory pathway of heart tissue. we applied echocardiographic techniques and transcriptomics analysis to detect changes in cardiac function indicators in sturgeon to further establish the link between heart tissue structure and pumping function under CHS. Meanwhile, the counteracting effect of aquaculture water cooling on CHS was systematically tested. Our research has established a foundation for the connection between heart structure and function in sturgeon under CHS and provides a reference for the protection of wild and farmed cold-water fish.

Materials And Methods

Fish maintenance

All animal handling procedures were approved by the animal care and use committee of Sichuan Agricultural University in accordance with the animal experiment guidelines under license NO. ZCY-2019202031. 400 healthy *A. baerii* of similar weights (84.531 ± 2.713 g) and body lengths (31.491 ± 1.746 cm) were purchased from Sichuan Runzhao Fishery Co., Ltd. China. The fish were held in tanks under a 12 h light : dark cycle with an uninterrupted oxygen supply to ensure dissolved oxygen remained above 7.5 mg/L. The water in the tanks was pretreated with UV light and an aeration process, and 20% of the culture water was renewed every day. The pH ranged between 7–8 and ammoniacal nitrogen and nitrite were maintained at 0–0.015 mg/L. The fish were fed with 1% of the fish body weight of commercial feed three times daily. Eventually, fish that were responsive, robust, and healthy were selected for experimentation.

Experimental treatment

According to the study by Yang et al., (in 2021) *A. baerii* exhibits significant increases in fatality and organ damage when the breeding water temperature reaches 28°C[14]. Based on the observations, the sturgeon in this experiment were divided among three groups to simulate normal and CHS conditions: a control group at 20°C (C) and two elevated temperature groups at 24°C (M) and 28°C (H). A total of 180 fish were randomly assigned to the three treatments, with each group consisting of four parallel tanks holding 15 fish each. For the control group, the sturgeon were raised at normal room temperature (20°C) for the duration of the experiment. The CHS groups were also held at room temperature for the first 14 days. Then, according to the heating scheme (Supplementary table 1), the temperatures in the stressed groups were increased by 1°C/d until reaching the experimental temperatures on day 22. The CHS temperatures were maintained until day 34 (heat stress exposure of 12 days). Immediately following, the water temperatures of the M and H groups were adjusted to 20°C for one week to allow for recovery (Supplementary table 1).

On day 35, 10 sturgeon were randomly selected from each group for sampling. Collect tail vein blood samples (n=5) for myocardial enzyme leakage detection, and cardiac tissue for transcriptomics (n=3) analysis. Fish from the recovery groups (R-M and R-M) were sampled as before on day 42.

Heart disease activity index (HDAI) evaluation

Based on the symptoms the sturgeon in the farm experienced in summer, we established a novel heart disease activity index (HDAI) with reference to the research by Chen et al. The index was designed to evaluate the "heart failure-like" behavior in *A. baerii* under CHS (Table 1) [38, 39]. Similar to the above experimental treatments (Section 2.2). The procedure was designed to assess the degree of damage to the *A. baerii* after experiment CHS. The monitored variables included feeding rate, swimming speed, imbalance rate, respiratory rate, cardiac color, and percentage of ventricular injury, and all variables were blind scored by two trained researchers. The evaluation experiment included CHS treatment and a recovery period.

Features graded	Description		
Feeding rate (n = 3)	0.7% - 1% (Containing 0.7%)	0	
	0.3% - 0.7% (Containing 0.3%)	1	
	0 - 0.3%	2	
Swimming speed (n = 6)	Normal (\leq 12 cm/s)	0	
	Mild fast (12 - 16 cm/s)	1	
	Speediness (\geq 16 cm/s)	2	
Imbalance rate	Less than 5% of the total number of experimental fish	0	
	5-15% of the total number of experimental fish	1	
	More than 15% of the total number of experimental fish	2	
Respiratory rate (n = 6)	Normal (\leq 70 /min)	0	
	Mild (70 - 90 /min)	1	
	Severe (\geq 90 /min)	2	
Cardiac color (n = 3)	Normal	0	
	Slightly black heart color		
	Severe darkening of the heart	2	
Percentage of ventricular injury (n = 3)	0		
	1 - 25%		
	26 - 50%		
	51 - 75%		
	76 - 100%	4	

Table 1 Scoring system of HDAI for sturgeon.

Feeding rate = total feed intake (g) / total body weight (g) $\times 0.01 \times 100\%$. To ensure measured swimming speed sturgeon are not subject to mechanical equipment and environmental impact. We used the camera to randomly record the swimming trajectory (curved or straight) and time of the horizontal sturgeon in the treatment tank one hour before feeding. Swimming speed = Swimming distance (cm) / time (s). Imbalance rate = number of lateral flip / total number. Meanwhile, the sturgeon was recorded for 5 minutes continuously one hour before feeding during the experiment. Six sturgeons in each group were randomly selected for respiration frequency statistics. Respiratory rate (T/min) = number of operculum

opening and closing cycles / time (min). Sensory score for the color of the dissected sturgeon heart (n = 3), then the hearts were dissected from sturgeon in each treatment and fixed in 4% paraformaldehyde overnight at room temperature. Paraffin sections were prepared and stained with haematoxylin and eosin (H & E). Sections were photographed under a light microscope (Nikon, Tokyo, Japan). The obtained heart histopathological images were analyzed using Image J 1.35a (National Institutes of Health, USA) image analysis software. Percentage of ventricular injury = ventricular injury area / total cross-sectional area of ventricle × 100%. Scoring system of HDAI for sturgeon can be seen in Table 1 (n \ge 3).

Myocardial enzyme determination

Myocardial enzyme is a biomarker of myocardial injury used to monitor the degree of myocardial necrosis [40, 41]. To further determine whether myocardial cells were necrotic, the content of myocardial enzymes in the plasma was measured. The myocardial enzyme leakage activity indexes for plasma were determined using diagnostic kits produced by Nan Jing Jian Cheng Bioengineering Institute (Nanjing, China) following the manufacturer's instructions. The blood from each sample was centrifuged at 3500 g 10 min at 4°C. The diagnostic kits used in this experiment measured LDH (lactate dehydrogenase) (A020-2-2), AST (aspartate aminotransferase) (C010-2-1), and CK (creatine kinase) (A032-1-1).

Echocardiography and histopathological examination

A modified version of the echocardiographic detection methods of Sun et al., Fang et al., and Liu et al. was used [42-44]. The sturgeon conducts the same time and treatment experiment as in section 2.2. Five sturgeon from each group (day 35 and 42) were randomly selected from the surviving pool of fish for echocardiographic analysis. Each selected fish was anaesthetized in 100 mg/L MS-222 (Sigma, USA) until loss of equilibrium, and placed ventral side up in a small sponge holder submerged in a transparent plastic bracket filled with water (water temperature corresponds to treatment temperature) from their respective aquaria. The ultrasound transducer (CHISON, China) was positioned on the ventral side of the sturgeon. Both the longitudinal and short axes of the CHISON Qbit 5VET Ultrasonic Diagnostic System (CHISON, China) were used during imaging. The frequency for B-mode imaging (Brightness-mode Ultrasound imaging) was 10 MHz and a typical sample of B-mode video of sturgeon has been provided in Supplementary Video 1. The frequencies of color Doppler and pulsed-wave Doppler were 6.5 MHz, presented in Supplementary Video 2. Assessments were performed so that the operators were 'blinded' to eliminate the potential for operator bias. Assessments were conducted in conditions temperature corresponding to the treatment temperature of the sturgeon for < 5 min, allowing echocardiographic recordings of at least three cardiac cycles. The water environment during the detection process and the indoor temperature during the detection are the corresponding experimental treatment temperatures.

With the help of the color-Doppler mode, sampling windows of pulsed-wave Doppler were positioned at the atrioventricular valve (AV), bulboventricular valve (BV), abdominal aorta (AA), and between the two valves. Clear images were captured and the Cardiac Software Package (CHISON, China) was used to analyze the relevant parameters. The pulsed-wave Doppler velocity measurement was taken from a

position as parallel to the blood flow as possible. The angle was adjusted if necessary, but adjustments were limited to a maximum of 15 degrees.

Basic parameters, such as body weight (BW), carcass weight (CW), full length (FL), and body length (BL) were measured promptly after echocardiographic examinations. Ratios of ventricular and heart mass to BW or CW were chosen to assess changes in cardiac morphology. Dissected tissues were immediately fixed in 10% neutral-buffered formalin for histopathologic examination. The samples were trimmed, dehydrated, and embedded in paraffin wax before sectioning at 4 µm for H & E and Masson trichrome staining. Images were recorded on a light microscope (Nikon, Tokyo, Japan). The degrees of change in fibrillar collagen content, atrophy, infiltration, necrosis, hyperplasia of fat, and blurred epicardial border in the heart were evaluated according to Huang et al [45]. Every change was assessed with a score (S) ranging from 0 to 6 depending on the degree of change: (0) unchanged, (2) mild change, (4) moderate change, and (6) severe change (diffuse lesion).

The obtained histopathological images were analyzed using Image J 1.35a (National Institutes of Health, USA) image analysis software. The basic calculation was conducted as follows: ventricle stratum spongiosum myocardial density (%) = pixels (area of myocardium) / pixels (total area of ventricle) ×100%; fibrillar collagen content in the myocardium of the stratum spongiosum = pixels (area of myocardial collagen) / pixels (total area of ventricle stratum spongiosum) ×100%.

Echocardiographic image data analysis

Clear echocardiographic images were obtained and offline analyses were performed. With reference to the analysis process established by Fang et al., [42], the measurement results were analyzed as follows. The epicardial border was outlined using the Cardiac Function Measurement Package (CHISON, China) and the end-diastolic (E-d) and end-systolic (E-s) ventricular areas (A) were calculated (Fig. 1). The diameter of the AV valve annulus during systole (D) was also recorded. In addition, ventricular filling (atrioventricular flow) and ejection flow (BV flow) were also measured (Fig. 1). Corresponding parameters such as peak velocity, mean velocity, and velocity time integral (VTI) can be automatically calculated when using the continuous wave Doppler mode. The calculated formulas-cross-sectional area (CSA) was used to evaluate stroke volume (SV, μ L), cardiac output (CO = SV*HR), and pulsatility index (PI = ventricular blood flow peak systole / end-diastolic trough). SV was calculated as CSA × VTI. Defining CSA as πD^2 / 4, the VTI of the BV valve was measured using pulsed Doppler. The pulsed Doppler image was used to estimate ventricular ejection time (VET) by measuring the time of ventricle ejection from start to finish (Fig. 1).

RNA-seq

2.7.1 RNA extraction, library preparation and illumina sequencing

The heart in sturgeon in different treatment group were selected for RNA-seq, each group including three replicates. Total RNA was extracted from the tissue using TRIzol® Reagent according the manufacturer's instructions(Invitrogen). Then RNA quality was determined using 2100 Bioanalyser (Agilent) and quantified using the ND-2000 (NanoDrop Technologies). High-quality RNA sample (OD260/280=1.8~2.2, OD260/230≥2.0, RIN≥6.5, 28S:18S≥1.0, >10µg) was used to construct sequencing library.

RNA-seq transcriptome libraries were prepared following TruSeqTM RNA sample preparation Kit from Illumina (San Diego, CA), using 1µg of total RNA. Shortly, messenger RNA was isolated with polyA selection by oligo(dT) beads and fragmented using fragmentation buffer. cDNA synthesis, end repair, A-base addition and ligation of the Illumina-indexed adaptors were performed according to Illumina's protocol. Libraries were then size selected for cDNA target fragments of 200–300 bp on 2% Low Range Ultra Agarose followed by PCR amplified using Phusion DNA polymerase (NEB) for 15 PCR cycles. After quantified by TBS380, Paired-end libraries were sequenced by Illumina NovaSeq 6000 sequencing (150bp*2, Shanghai BIOZERON Co., Ltd).

2.7.2 De novo assembly and annotation

The raw paired end reads were trimmed and quality controlled by Trimmomatic with parameters (SLIDINGWINDOW:4:15 MINLEN:75) (version 0.36

http://www.usadellab.org/cms/uploads/supplementary/Trimmomatic).Then clean data from all samples were used to do RNA de novo assembly with Trinity (http://trinityrnaseq.sourceforge.net/). All the assembled transcripts were searched against the NCBI protein nonredundant (NR), String, and KEGG databases using BLASTX to identify the proteins that had the highest sequence similarity with the given transcripts to retrieve their function annotations and a typical cut-off E-values less than 1.0×10⁻⁵ was set. BLAST2GO (http://www.blast2go.com/b2ghome) program was used to get GO annotations of unique assembled transcripts for describing biological processes, molecular functions and cellular components. Metabolic pathway analysis was performed using the Kyoto Encyclopedia of Genes and Genomes (KEGG, http://www.genome.jp/kegg/).

2.7.3 Differential expression analysis and functional enrichment

To identify DEGs (differential expression genes) between two different samples, the expression level of each transcript was calculated. Fragments/KB/Million reads (FPKM) were used to analyze the expression levels of differential genes. RSEM (http://deweylab.biostat.wisc.edu/rsem/) was used to quantify gene and isoform abundances. R statistical package software EdgeR (Empirical analysis of Digital Gene Expression in R, http://www.bioconductor.org/packages/2.12/bioc/html/edgeR.html) was utilized for differential expression analysis. Meanwhlie, functional-enrichment analysis including GO and KEGG were performed to identify which DEGs were significantly enriched in GO terms and metabolic pathways at Bonferroni-corrected P-value ≤0.05 compared with the whole-transcriptome background. GO functional enrichment and KEGG pathway analysis were carried out by Goatools (https://github.com/tanghaibao/Goatools) and KOBAS (http://kobas.cbi.pku.edu.cn/home.do).

Statistical analysis

Continuous variables were expressed as mean \pm standard deviation (SD). Categorical variables were expressed as counts and percentages. Data were analyzed using SPSS version 27. Differences between groups were assessed with one-way ANOVAs followed by post hoc analyses with Dunnett's tests. Differences between the means were considered significant or extremely significant at the 95% or 99% confidence levels (*P* < 0.05 or *P* < 0.01), respectively.

Results

Systemic symptoms of heart failure in A. baerii caused by CHS

A. baerii developed relatively obvious heart failure symptoms after CHS treatment. Starting from day 1 to 19, the HDAI scores of the fish in group H were significantly (P < 0.05 or P < 0.01) higher than those of the control group (Fig. 2A). Notably, the cardiac HDAI in group H abruptly peaked from day 4 to day 12, during which time it remained at a high level. Specifically, there were significant increases in feeding rate, swimming speed, imbalance rate, respiratory rate, cardiac color, and percentage of ventricular injury scores relative to the control group for most of the days between 4–12 (Fig. 2C). Fish in the heat-stressed group showed a significant recovery trend from day 18 to 19 (P < 0.05). The survival rates of the group H was significant lower than the control group (Fig. 2B). The survival rate dropped significantly from day 5 and continued to decrease until the end of the heat stress period. During the recovery period, there was a clear recovery trend in the M group, while the survival rate of the H group continued to decline (Fig. 2B). Generally, sturgeon in the heat stress groups showed systemic cardiopathic symptoms. In the control group, almost no deaths occurred during the experiment, no obvious pathological symptoms were observed, and the HDAI scores remained relatively stable without significant change.

Significant irregular ventricular shape and heart atrophy in A. baerii caused by CHS

Immediately after the CHS and following recovery, we performed further measurements of morphological indicators of heart condition and basic parameters of the health of *A. baerii*. Severe abnormal shapes, darkening of the ventricle, and atrophy of the heart increased with the intensity of CHS, but were partially relieved after the recovery period (Fig. 3A). Specifically, the ventricle darkening and ventricle atrophy were the most obvious with increasing CHS intensity. Next, the basic parameters were measured. All morphological indicators of the heart showed a decreasing trend with increasing intensity of CHS. The indexes related to heart weight showed significant reductions in group H, but returned to normal levels after the recovery period. Meanwhile, there was a tendency for the ventricles to atrophy under CHS (Table 2).

Table 2 Morphological changes of *A. baerii* hearts after CHS and recovery..

ltem	C	M	Н	R-M	R-H
BW, g	86.562 ±	86.769 ±	84.692 ±	90.671 ±	90.589 ±
	19.520	19.391	15.432	15.889	12.751
CW, g	78.298 ±	78.177 ±	77.292 ±	82.208 ±	79.749 ±
	17.705	16.924	14.465	15.021	12.555
TL, cm	32.308 ±	32.485 ±	31.954 ±	33.708 ±	33.208 ±
	1.645 ^{ab}	1.743 ^{ab}	1.961ª	1.850 ^b	1.605 ^{ab}
BL, cm	25.427 ± 1.459	25.631 ± 1.478	25.454 ± 1.616	25.715 ± 1.208	25.400 ± 1.238
HM, g	0.152 ± 0.042^{ab}	0.164 ± 0.051 ^a	0.127 ± 0.029 ^b	0.157 ± 0.039 ^{ab}	0.170 ± 0.057 ^{ac}
VM, g	0.0710 ±	0.0782 ±	0.0590 ±	0.0702 ±	0.0808 ±
	0.0205	0.0324	0.0124	0.0164	0.0247
HM /	0.1938 ±	0.2078 ±	0.1661 ±	0.1926 ±	0.1991 ±
CW	0.0301 ^{ab}	0.0336 ^a	0.0288 ^b	0.0336 ^{ab}	0.0907 ^{ac}
VM /	0.1080 ±	0.1041 ±	0.0879 ±	0.0953 ±	0.0957 ± 0.012
CW	0.0236	0.0272	0.009	0.0196	
RHM	0.1752 ±	0.1874 ±	0.1513 ±	0.1729 ±	0.1752 ±
	0.0268 ^{ab}	0.0298 ^a	0.0259 ^b	0.0307 ^{ab}	0.0813 ^{ac}
RVM	0.0966 ±	0.0938 ±	0.0798 ±	0.0851 ±	0.0859 ±
	0.0211	0.0239	0.0087	0.0177	0.0107

Values are mean \pm SD, n = 5–13 hearts for each groups. BW: Body weight; CW: Carcass weight; TL: Total length; BL: Body length; HM: Heart mass; VM: Ventricular mass; RHM: Relative heart mass = [(HM / BW) × 100]; RVM: Relative ventricular mass = [(VM / BW) × 100]. a, b and c represent significantly different groups (P < 0.05).

Fibrillar collagen level increased and cardiomyocyte necrosis caused structural damage to the heart

To observe the histopathological changes of *A. baerii* heart failure, heart sections were stained following the protocols of H & E and Masson trichrome staining. Masson's trichrome staining showed that the changes were extremely pronounced in the compact layer in the heat-stressed group (Fig. 3A). The most obvious changes were the decrease in the thickness of the compact layer and the dramatic increase in the content of fibrillar collagen (Fig. 3A, B and E). Meanwhile, CHS also caused the density of the spongy layer to significantly decrease and the fibrillar collagen content to significantly increase (Fig. 3C, D). All indices showed different degrees of recovery after the recovery period (Fig. 3).

Sections were systematically examined, revealing that the most obvious lesions were mainly concentrated in the ventricular tissue. Specifically, H & E staining showed massive necrosis of cardiomyocytes in the compact layer of the ventricle in group H, exhibiting symptoms of myocardial atrophy and inflammatory cell infiltration (Fig. 4A c, f, g). The histopathological score of the compact layer also significantly increased (P < 0.05) (Fig. 4A h). CHS also resulted in significant increases in epicardial fat and increased blurring of the boundary between the compact layer and the epicardium compared to the control (Fig. 4A g). To further confirm the presence of necrosis in the cardiomyocytes we tested the myocardial enzyme leakage levels in the plasma. In general, the levels of myocardial enzymes exhibited an upward trend under CHS, but the increase in CK was most obvious (Fig. 4D) (P < 0.05).

The regulatory mechanism of cardiac fibrosis and myocardial necrosis under CHS

To study the specific role of cardiac fibrosis and myocardial necrosis in the heart of *A. baerii*, we selected the group C, H and R-H heart for transcriptome sequencing. Transcriptomics analysis showed that fibrous collagen deposition increased after CHS. It is manifested as an increase in the production of collagen fibers (*COL1A, COL2A, COL4A* and *COL6A*) and a down-regulation of the expression of genes that regulate the degradation of collagen fibers (*COLase* 3 and *COLase* 4). In the necroptosis regulatory network, down-regulated of the *Caspase* 8, leading to inhibition of apoptotic pathways in cardiomyocytes and shifting to necrosis. The present studies have shown that mitochondrial fission may be an important inducement of cardiomyocyte necrosis, accompanied by the significant up-regulation of *DNM1L* and *Hsp* 90 after CHS (Fig. 6L and M). However, timely cooling can alleviate the process of cardiac fibrosis and cardiomyocyte necrosis to a certain extent.

Blocked heart activity under severe CHS

To further investigate the association between changes in cardiac tissue structure and cardiac function, echocardiography combined with B-mode imaging and Doppler imaging were used. In the current study, both the ventricular end-diastolic area (VEDA) and ventricular end-systolic area (VESA) showed increasing trends in CHS conditions, with significant differences in the VESA (P < 0.05). After the recovery period, the M group remained unchanged, while the VEDA and VESA continued to rise in the H group (P < 0.05) (Fig. 7A and B). CHS resulted in a significant reduction in atrioventricular valve diameter, which did not return to normal levels after the recovery period (Fig. 7C). Meanwhile, a marked HR increase was observed in the H group compared to control group, but after recovery the HR was no longer elevated (Fig. 7D).

CHS caused decrease in heart ejection effectiveness

Impaired cardiac activity may directly change blood flow parameters. Therefore, ultrasound Doppler analysis was used to analyze the changes in cardiac blood flow velocity. After CHS, there was no significant change in AA blood flow velocity in group M, but group H showed a significant increase (Fig. 8). The blood flow velocities from the BA and AV were significantly reduced in the recovered fish from group H compared to the control fish (Fig. 8B, C) (P < 0.05). Except group M and H, no significant differences in AA blood flow velocity were detected among groups.

In addition, indicators of cardiac function were measured using the Cardiac Function Measurement Package. With increasing CHS intensity, SV, CO, and VET showed significant and sequential decreases in the M and H groups compared to the control group (P < 0.05) (Fig. 7F, G and H). However, the opposite trend was true for the PI of group H (Fig. 7E). Considering the trends in HR, which suggested decreased myocardial contractility and increased cardiac afterload, the changes could be characterized as those of a typical high-resistance and low-output heart failure. During the recovery period, almost all indicators tended to return to normal levels (group C), in particular, SV, CO, and VET were significantly increased in the R-H group (P < 0.05) (Fig. 7F, G and H). However, the SV and CO levels did not return to normal levels in group H.

Discussion

CHS caused by the increase of water temperature in aquaculture ponds in summer may lead to heart failure of sturgeon and affect the development of sturgeon industry. However, how CHS affects the changes in sturgeon's heart structure and function is still unclear and its effect on fish production significance for aquaculture need to be further studied. This study explored the changes in the structure and function of the sturgeon's heart under CHS, and tried to establish the connection. Combined with histopathology, echocardiographic detection technology and transcriptomics analysis, it was found that CHS caused *A. baerii*'s ventricular fiber collagen content to increase and myocardial necrosis, which caused *A. baerii*'s cardiac dysfunction and decreased pumping effectiveness. After the recovery period, although most results showed that the heart function of the heat-stressed group had a tendency to recover. However, there were still injuries in group H that could not eliminate the symptoms. It indicates that the sturgeon heart has a certain ability to regulate itself, which may be accomplished through the regulation of fiber collagen content and the proliferation of myocardial cells [35, 46, 47].

The present study identified the proliferation of myocardial interstitial myocardial collagen and myocardial necrosis as important factors leading to the cardiac imbalance and heart failure. Myocardial fibrosis is an important causes of cardiac dysfunction in *A. baerii*. Myocardial fibrosis is a pathological reaction that causes an imbalance in the collagen ratio due to increased deposition of collagen in the myocardial interstitium, which may induce a variety of cardiovascular diseases [26, 48]. Studies have shown that increased ventricular stiffness and decreased cardiac function have been observed in Atlantic cod (*Gadus morhua*) at warm acclimation [31]. Meanwhile, in zebrafish, where there is significantly less thick collagen fibres in the hearts of fish acclimated to 20°C compared with those acclimated to 28°C [46]. We guess that the zebrafish myocardial collagen fibres will show the opposite trend after warm acclimation. Fibrous collagen deposition increases ventricular stiffness. Similarly, after CHS, the proportion of *A. baerii*'s ventricular fiber collagen increased significantly in group H. Fibrous collagen increases the passive stiffness of the chamber wall, and excessive myocardial fibrosis will reduce chamber compliance and dilatability, which may have a significant impact on ventricular diastolic filling

[28]. We believe that it is caused by delayed or hindered degradation of collagen fibers and over-synthesis to maintain part of the heart function. Therefore, we observed that the VEDA of the ventricle remains unchanged but the VESA increases significantly. Meanwhile, CO and SV in the CHS group were significantly reduced and implying decreased pumping effectiveness. Such changes are common, and permanent, in the hearts of patients suffering from cardiac hypertension, dilated cardiomyopathy or chronic congestive heart failure, and collagen fiber increased greatly contribute to the associated diastolic dysfunction and eventual heart failure [49-52]. Considering the importance of the ventricles to the whole heart, this may be an important factor in inducing heart failure in sturgeon. Study found that the cold-induced increase in salmon heart collagen deposition was reversed after experiencing chronic warming [23, 35]. Therefore, the significant decrease in collagen content of *A. baerii*'s ventricular fibers during the recovery period can be attributed to the degradation of collagen.

Furthermore, cardiac myocytes necrosis, which is one of the main pathological changes in the hearts of A. baerii under CHS, can cause cardiac atrophy and dysfunction [32, 53]. Study found that the H₂O₂ consumption capacity of cardiac mitochondria in rainbow trout decreased with increasing temperature, which may increase the risk of oxidative stress and cause cardiomyocyte necrosis [54]. In present study, histopathology and myocardial enzyme levels confirmed the presence of necrosis of myocardial cells. And transcriptomics studies have found that cardiac mitochondrial damage may mediate myocardial cell necrosis. Experiment found that the delayed delivery of ion channel flux action potentials caused by the collapse of cardiac myocytes may seriously affect pumping in fish hearts. [55]. However, as an increase in physiological temperature increases HR in fish [16, 56], this suggests that the heart is pumping less blood per beat at a faster rate. Therefore, we have observed an increasing trend of the AA flow velocity under chronic heat stress (group H) in sturgeon, but the BV and AV flow do not change significantly. It explains the significant increase in A. baerii heart rate and the significant decrease in SV after CHS (group H). Meanwhile, damage to myocardial structure will cause insufficient energy supply, abnormal ventricular contraction (increased ventricular volume at the end-systolic area), and lead to a decrease in stroke volume and pumping efficiency [57]. Therefore, cardiac function in A. baerii under heat stress is imply a reduction in myocardial contractility and an augmentation of cardiac afterload, matching the typical characteristics of high-resistance and low-output heart failure.

Like most cold-water fish fish, the sturgeon's living space and life activities are under threat and being compressed by rising water temperatures, which is apparent in the Yangtze River Basin [4, 6]. In addition, sturgeon grow slowly and usually take 7–8 years to reach sexual maturity [20]. Cardiac dysfunction is undoubtedly an important risk factor that could hinder the growth and development of sturgeon, potentially increasing their age at maturation. Due to the inability of their hearts to quickly respond to high water temperatures, most sturgeon could exhibit "cardiac syndrome" (like *S. salar*) in the high water temperature environment in summer [58, 59]. According to the results of the echocardiography and microstructure observations after the recovery period, it was found that the sturgeon's heart was unable to completely return to normal working levels. Current evidence suggests that the upper limits of temperature tolerance in fish are primarily determined by the points at which their hearts fail to maintain

tissue perfusion and adequate oxygenation [60, 61]. Disorders of heart function may affect fish swimming performance, predator avoidance, foraging behavior, and alter a species geographic distribution [62, 63], leading to the local extinction of fish species [4]. Therefore, avoiding high water temperature sturgeon farming and seeking effective cooling methods may be important directions for sturgeon farming in the future. In addition, behavioral observations of juvenile sturgeon should be performed regularly during the conservation and breeding of endangered sturgeon species, and antistress prevention products should be added appropriately to improve heart health.

Conclusions

In conclusion, we have confirmed that timely cooling of the culture water can alleviate the heart failure of *A. baerii* caused by CHS. CHS can cause *A. baerii* to show obvious "heart failure" characteristics, and cardiac function examination reveal a reduction in myocardial contractility and an augmentation of cardiac afterload, matching the typical characteristics of high-resistance and low-output heart failure characteristics. The rate of *A. baerii* 's cardiac fibrosis and myocardial necrosis was relieved after the culture water was cooled. Therefore, the present study shows that maintaining normal growth water temperature and timely cooling during cold-water fish farming are important measures for healthy farming.

Declarations

Ethics approval statement

All animal handling procedures were approved by the animal care and use committee of Sichuan Agricultural University in accordance with the animal experiment guidelines under license NO. ZCY-2019202031.

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Authors' contributions

All authors read and approved the final manuscript. Chaoyang Zhang, Miao Zhang and Zihan Xu contributed the work equally. All authors read and approved the final manuscript. Chaoyang Zhang: Writing original draft, Methodology, Investigation. Miao Zhang: Investigation. Zihan Xu: Data curation, Software. Datian Li: Data curation. Quan Gong: Methodology, Software. Xiaoli Huang: Formal analysis, Resources, Validation. Jiayun Wu: Formal analysis. Zhi He and Xiaogang Du: Data curation, Supervision, Investigation. Defang Chen and Jun Jiang: Supervision, Investigation. Jun Du: Resources and Formal analysis. Shiyong Yang: Project administration, Writing original draft, Investigation, Methodology, Conceptualization, Contribution.

Declaration of Competing Interest

Authors declare that the research was conducted without any commercial or financial relationships that could be construed as a potential conflict of interest.

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Availability of data and materials

All data generated or analyzed were included in the article. All materials are available from the corresponding author, on reasonable request.

Ethics approval and consent to participate

All animal handling procedures were approved by the animal care and use committee of Sichuan Agricultural University in accordance with the animal experiment guidelines under license NO. ZCY-2019202031.

Consent for publication

Not applicable.

Competing interests

The authors declare no conflict of interest.

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Figures



Figure 1

The echocardiography detection procedures for analysis of various cardiac parameters in A. baerii. Typical echocardiographic images of the sturgeon, showing the epicardial boundary, ventricular filling



flow (AV flow), and ventricular ejection flow (BV flow). Relevant parameter values were automatically generated by the Cardiac Function Measurement Package.

Figure 2

The HDAI (A) and survival rate % (B) of A. baerii in different temperature groups at different time points. HDAI component parts, including: feeding rate; swimming speed; imbalance rate; respiratory rate; cardiac color; and percentage of ventricular injury (C). Survival curves were estimated by the Kaplan-Meier method, and significance analysis was performed by the log-rank test. P < 0.05 (*) and P < 0.01 (**) indicate that groups are significant or extremely significant different from the control group.



Figure 3

Morphologies, thicknesses and fibrillar collagen content of A. baerii hearts or ventricles in the control, after CHS and the recovery period. A: The dynamic process of cardiac morphology and the histopathology of the ventricle (Arrowheads indicate ventricular compact layer thickness, dashed lines highlight myocardial collagen and red arrows indicate black areas in the ventricles and arterial bulbs. V: Ventricle; A: Atrium; BA: Bulbus arteriosus.) B: Thickness of the compact layer; C: Density of the sponge

layer; D and E: Content of fibrillar collagen in the myocardium of the sponge and the cavernous layer, respectively. Different lowercase letters indicate significant differences between groups (P < 0.05).



Figure 4

Histopathological examination of the ventricles and myocardial enzymes detection in the plasma of A. baerii under CHS. A: Histopathological examination of the ventricles of A. baerii. a: Healthy heart of A. baerii; b: Massive infiltration by inflammatory cells in the spongy layer (Group M); c and f: Ventricular

spongy myocardial necrosis (Group H); d and g: Infiltration of inflammatory cells in the ventricle, blurring and collagenization of the epicardial border (Group H); e: Ventricular microvascular injury in Group H (\rightarrow : necrotic cells; \rightarrow : nucleolysis; \rightarrow : karyopyknosis; \blacksquare : inflammatory cell infiltration; \bigcirc : myofibrillar atrophy of the dense layer of the ventricle; \blacklozenge : myocardial atrophy in the dense layer; \blacktriangle :blurred epicardial border; triangular arrowhead: pericardial myocardial collagen); h: Overall health status (total index) of ventricles in different groups based on histopathological lesions. A: LDH; B: AST; C: CK. Different lowercase letters indicate significant differences between groups (P < 0.05).

Figure 5

DEGs of fibrous collagen regulation-axis in A. baerii. Mapping the fibrous collagen regulation-axis network in the heart of A. baerii based on transcriptome data. A-I: the column map showed the expression level of reads of regulate collagen deposition genes in the transcriptome (n = 3, data were presented as means \pm standard deviation). * or *** represents a significant or highly significant difference (P < 0.05 or P < 0.001) between two groups.



Figure 6

DEGs of necroptosis regulatory pathway in A. baerii. A: Mapping the necroptosis regulatory network in the heart of A. baerii based on transcriptome data. The red and green character indicate the up regulation and down regulation of DEGs, respectively. B-K: the column map showed the expression level of reads of regulate necroptosis regulatory pathway genes in the transcriptome (n = 3, data were presented as means



 \pm standard deviation). *, ** or *** represents a significant, extremely or highly significant difference (P < 0.05, P < 0.01 or P < 0.001) between two groups.

Figure 7

Ventricular activity, outflow and inflow parameters in A. baerii measured by B-mode and Doppler echocardiography. A and B: Ventricular end-diastolic area (VEDA) and ventricular end-systolic area (VESA), respectively, evaluated by area (mm2); C: Atrioventricular annulus diameter; D: Heart rate. E: PI; F: SV; G: CO; H: VET. All echocardiographic results were standardized by body weight. Different lowercase letters indicate significant differences between groups (P < 0.05).



Figure 8

Ventricular activity, outflow and inflow parameters in A. baerii measured by B-mode and Doppler echocardiography. A and B: Ventricular end-diastolic area (VEDA) and ventricular end-systolic area (VESA), respectively, evaluated by area (mm2); C: Atrioventricular annulus diameter; D: Heart rate. E: PI; F: SV; G: CO; H: VET. All echocardiographic results were standardized by body weight. Different lowercase letters indicate significant differences between groups (P < 0.05).

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