



The effects of increased temperature and CO₂ on the Amazon fish, tambaqui (*Collosoma macropomum*, Cuvier, 1818)

Amtyaz *¹, Ramon Barros Baptista ², Vera Maria Fonseca de Almeida e Val ^{2&3}

¹ Dept. of Zoology, Sir Syed Govt. Girls College Nazimabad, Karachi, PAKISTAN.

² Laboratory of Eco-Physiology and Molecular Evolution (LEEM), National Institute for Research in the Amazon, INPA. Manaus, Brazil

³ Laboratory for Applied Genetics in Aquaculture (AGA), University Nilton Lins (UNL), Manaus, AM, Brazil

*Corresponding author E-mail: intiazsafi76@gmail.com

Abstract

The effects of temperature and carbon dioxide (CO₂) changes on different haematological parameters and enzymes activities such as LDH (Lactate de Hydrogenase), MDH (Malate de Hydrogenase) & GST (Glutathione-S-Transferase) were studied in the liver and skeletal muscles of Tambaqui fish (*Collosoma macropomum*, Cuvier, 1818) in the Laboratory of Ecophysiology and Molecular Evolution (LEEM) of the National Institute for Amazonian Research (INPA). The experimental procedures involved the exposure of fish to temperature rise from 28C° to 32C° and an increased in CO₂ concentration up to 10 times higher than the current levels for 24, 48, 72 and 96 hours. The levels of LDH, GST and MDH in both the tissues of Tambaqui (Muscles & Liver) showed a significant decreased at different along the different exposure periods. The levels of plasma glucose in blood decreased and the values of Hb, Ht, RBC, VCM, and HCM & CHCM showed increased after the exposure.

Keywords: Temperature, CO₂, Tambaqui fish, Amazon, Brazil.

1 Introduction

Owning one of the greatest biological diversities of the world, the most notable fauna of the Amazon basin is the fish one, whose numbers of species are estimated to be more than 2500. Among the water bodies of the region, the biggest variety of fish is found in Amazon River, presenting a total of 256 species. This number is extremely high compared to the whole Europe hydrographic basins, where only 192 species are registered [1]. The adaptations of fish to the Amazonian flooding areas include several strategies concerning their growth and reproduction. However, they are also characterized for presenting complex morphological, anatomical, physiological, and behavioral adaptations to ensure the survival in an ever-changing environment [2]. A great diversity of species has corresponds to a wide range of biochemical, physiological, phenological and ethological strategies, as well as a range of anatomical and morphological variations to deal with periodical flooding, making the studies in floodable areas very complex. However, an innovative approach may enable the detection of a mechanism for common regulation to all floodable biota components, with benefits for the scientific knowledge, for the productive chain and especially to people in the region.

The Amazon unquestionably presents the largest fresh water resource and fish diversity in the world. In this region, fish is the most important dietary protein source to its population (Per capita consumption of 60 kg/year) and fishing is among the most important economic activity, involving more than 110,000 people [3], [4].

Tambaqui (*Colossoma macropomum*) is the most important commercial species in the Amazon and the main Brazilian native species for aquaculture. A large number of studies on its culture have been made available in recent years, emphasizing its performance and tolerance to intensive culturing conditions ([5], [6], [7], [8], [9], [3], [10], [11], [12], [13], [14]). However, information on the effect of Temperature and CO₂ on Tambaqui is still scarce and required.

The tambaqui (*Colossoma macropomum*) is a freshwater fish of the subfamily; Serrasalminae, family; Characidae and is the largest characin of South America. It is also known by the names pacu, black pacu, black-finned pacu, giant pacu, cachama and gamitana. The tambaqui is found in the Amazon and Orinoco basins in its wild form. It may reach more than 1 meter in total length and 30 kilograms in total weight. It is similar in shape to the Piranha and is

sometimes confused with the carnivorous fish. It is tall and laterally compressed with large eyes and a slightly arched back. Body color is basic black to gray with spots and blemishes in its mid body. All the fins are black and the pectoral fins are small. Around 10 percent of a tambaqui's weight is fat. This species is usually solitary. Adults stay in flooded forests during the first 5 months of flooding and consume fruits and grains. Young and juveniles live in black waters of flood plains until sexual maturity. The tambaqui feeds on zooplankton, insects, snails, and decaying plants. Research has indicated that this species plays an important role in dispersing seeds from fruits. The tambaqui is used in aquaculture because it can live in mineral poor waters and is very resistant to diseases. This species is marketed fresh and frozen. The aim of the present study was to evaluate the effect of water exchange regime as an alternative to minimize aquaculture environmental impact using tambaqui physiological parameters and performance as indicators.

Temperature and CO₂ are the major environmental factors, and play a critical role in growth, reproduction, immigration, succession and metabolism and health of ecosystems. A rise in temperature and CO₂ changes the physical and chemical properties of water. High water temperature induce increased metabolic rate in the aquatic organisms, resulting for example, in more demand for food in fishes. . In recent years, there was an increase of interventions in the Amazon - such as new hydroelectric plants, changes in the soil, mining and roads - all of which had immediate impacts on the aquatic environment. As this fish is of commercial interest, it is very important for the Amazon, so it is important to know what reactions will be caused by climate changes on Tambaqui. The responses of Tambaqui fish (*Collossoma macropomum*, Cuvier, 1818) to the stresses associated with acidification and high temperature is still not clearly understood, and a number of recent experiments have addressed this problem. The current work attended to expose tambaqui to higher temperature and higher CO₂ to check the main responses to the acute hypercapnia in high temperature.

In the present project we had studied the effect of two major environmental factors on Tambaqui fishes (*Collossoma macropomum*) of Manaus, Amazona's state of Brazil. We have, than, studied the effect of acute increased in temperature and CO₂ on LDH (Enzyme for an-Aerobic metabolism), MDH (Enzyme for Oxidative aerobic metabolism, present both in mitochondria and cytoplasm) and GST (an Anti-oxidant Enzyme, also responsive to xenobiotic exposure) from muscles and liver of Tambaqui fish (*Collossoma macropomum*, Cuvier, 1818).

2 Materials & methods

The experiment was performed in the Laboratory of Ecophysiology and Molecular Evolution (LEEM) of National Institute for Research in the Amazon, INPA of Amazonia state of Brazil, during the period between May and June of 2012. Tambaqui (*Collossoma macropomum*), were obtained from a fish farming station, located near Manaus, Amazonas State, Brazil, and fish were transported to LEEM Laboratory at INPA, and kept in tanks (1000 L) supplied with continuously water flow and aeration at 28C°, 12L: 12D photo period and fed for one month. After the acclimatization period, fish were divided into two groups: control (28C° normocapnia) and experimental (E: 32C° hypercapnia) groups (n = 6, each group) and assayed after 96h exposure. The control and experimental groups were transferred to the six aquaria (5L, one fish per aquarium) containing water with the same physical and chemical characteristics as those of the acclimatization period. Feeding was suspended 24h before the experiments. Biometry was made in the beginning of the experiment aiming to assess total body length in cm, standard body length in cm, body height in cm and body weight in g. As a result of the experiments we sampled three groups: C (Control 0h); CC (Control 96h) and E (Experimental 96h).

Fishes were killed by medullar section according to Brazilian recommendations of animal care council from the Brazilian Ministry of Science, Technology and Innovation (CONCEA – MCTI). Blood samples were drawn from the caudal vessel using heparinized syringes for determining blood tests before fish death. For the Enzymatic analysis (LDH, MDH and GST) liver and muscle were isolated, immediately frozen in liquid N₂, and stored at -80°C until processing. Tissues were washed in cold 250 mM sucrose solution and homogenized in cold 100 mM phosphate buffer (1:1, w/v), PH 7.0, with a Potter-Elvehjem homogenizer. The homogenates were centrifuged at 18,000g in a Sorvall RC-5B during 30 min at 4°C. For the haematological parameters evaluation, hematocrit values (Ht %) were determined after 10 minutes centrifugation in micro-haematocrit tubes. Haemoglobin concentration ((Hb) mg.dL⁻¹) was measured as described by Kampen & Zijlsta (1964). Red blood cell counts (RBC) were obtained with the use of a Neubaur chamber. Mean corpuscular volume (MCV), mean corpuscular haemoglobin concentration (MCHC), and mean corpuscular haemoglobin (MCH) were determined according to Brow (1976). The glucose levels were determined by electronic meter blood glucose, Accu-chek advantage II. For the enzymatic parameters evaluation, the activity of lactate dehydrogenase (LDH; E.C.1.1.1.27), Malate dehydrogenase (MDH; E.C. 1.1.1.37) and GST were estimated in skeletal muscle and liver according to Driedzic & Almeida-Val (1996). For analysis of LDH, MDH & GST activities, samples were homogenized in a buffer containing 150 mM Imidazole, 1mM EDTA, 5mM

dithiothreitol (DTT) and 1 % Triton X-100, PH7.4. To avoid loss of enzyme activity during sample preparation, procedures were performed on ice. Data is presented as Means and Standard Deviation of the Means and were compared.

Table.1: Change in different blood parameters of tambaqui (*Colossoma macropomum*) exposed to different temperature and CO₂.

	Glucose mg/dl	Abs 540 nm	[Hb] g/dl	Ht (%)	RBC (x10 ⁶ mm ³)	VCM	HCM	CHCM
C1-0h	28.00	0.112	3.27	19.00	1.39	136.69065	23.528058	17.212632
C2-0h	38.00	0.124	3.62	23.00	1.49	154.36242	24.300671	15.742609
C3-0h	33.00	0.144	4.20	24.00	1.51	158.9404	27.846358	17.52
C4-0h	44.00	0.126	3.68	23.00	1.5	153.33333	24.528	15.996522
C5-0h	33.00	0.151	4.41	25.50	1.55	164.51613	28.446452	17.29098
C6-0h	41.00	0.147	4.29	25.00	1.33	187.96992	32.273684	17.1696
Average	36.17		3.91	23.25	1.46	159.30	26.82	16.82
Std. Dev.	5.40		0.41	2.12	0.08	15.39	3.05	0.69
CC1-96h	15.00	0.162	4.73	27.00	1.21	223.1405	39.094215	17.52
CC2-96h	28.00	0.162	4.73	23.50	1.14	206.14035	41.494737	20.129362
CC3-96h	29.00	0.216	6.31	27.00	2.15	125.5814	29.335814	23.36
CC4-96h	66.00	0.209	6.10	30.50	1.61	189.44099	37.90559	20.00918
CC5-96h	56.00	0.222	6.48	27.00	1.77	152.54237	36.623729	24.008889
CC6-96h	15.00	0.178	5.20	19.50	1.27	153.54331	40.925984	26.654359
Average	34.83		5.59	25.75	1.53	175.06	37.56	21.95
Std. Dev.	7.804912983		0.73	3.45	0.36	33.92	4.04	3.03
E1-96h	40.00	0.21	6.13	28.00	1.58	177.21519	38.810127	21.9
E2-96h	30.00	0.242	7.07	28.00	2.35	119.14894	30.069787	25.237143
E3-96h	149.00	0.238	6.95	27.00	1.89	142.85714	36.77037	25.739259
E4-96h	23.00	0.219	6.39	28.50	1.83	155.7377	34.944262	22.437895
E5-96h	30.00	0.189	5.52	27.50	1.69	162.72	32.66	20.07
E6-96h	46.00	0.246	7.18	31.50	2.22	141.89189	32.356757	22.80381
Average	33.80		6.54	28.42	1.93	149.93	34.27	23.03
Std. Dev.	9.12		0.59	1.46	0.27	18.28	2.92	1.94

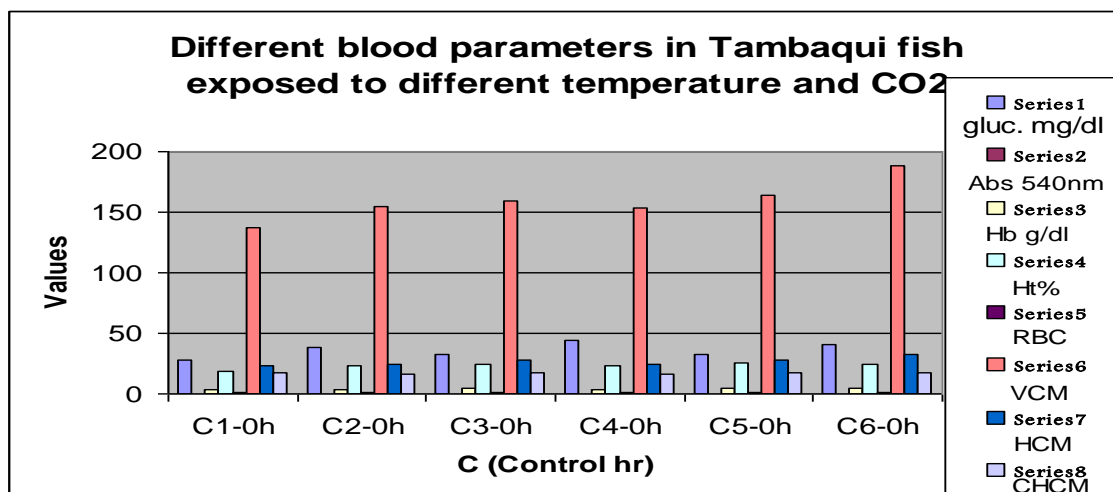


Fig. 1: Different blood parameters in C(Control 0hr)

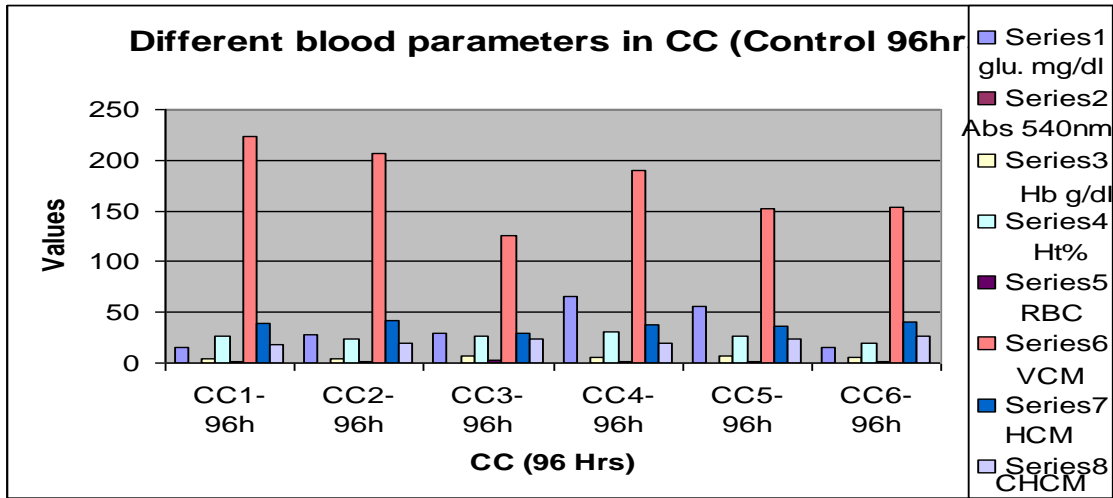


Fig. 2: Different blood parameters in CC(Control 96hrs)

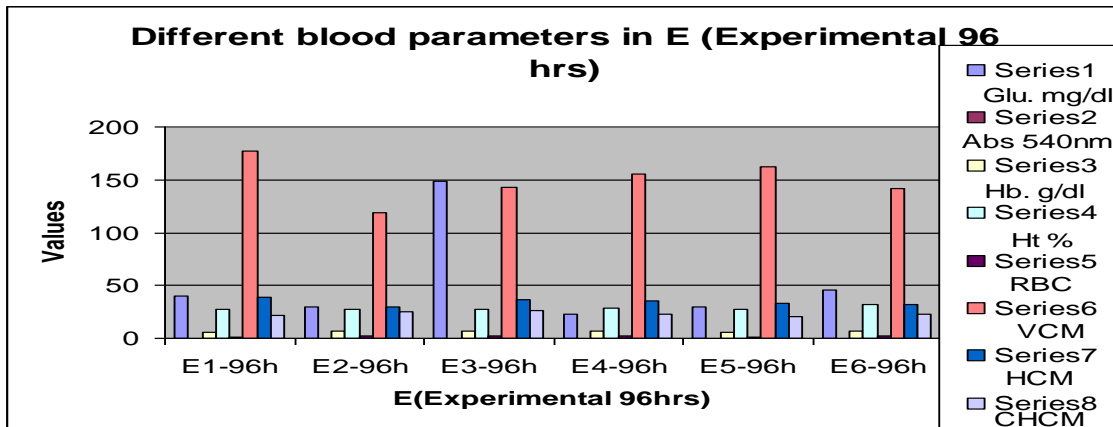


Fig. 3: Different blood parameters in E(Experimental 96hrs)

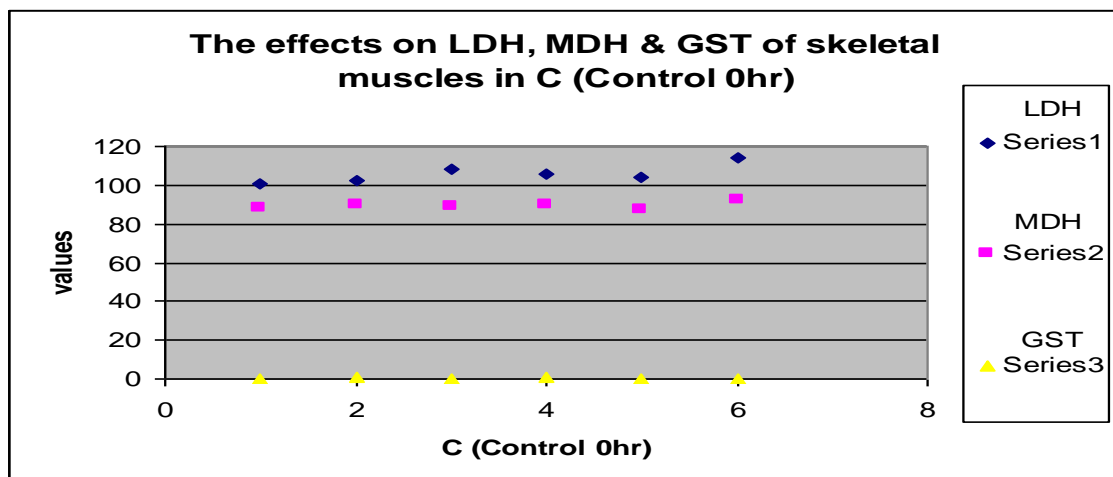


Fig. 4: The effects on LDH, MDH & GST of Skeletal muscles in C (Control 0hr) of Tambaqui fish.

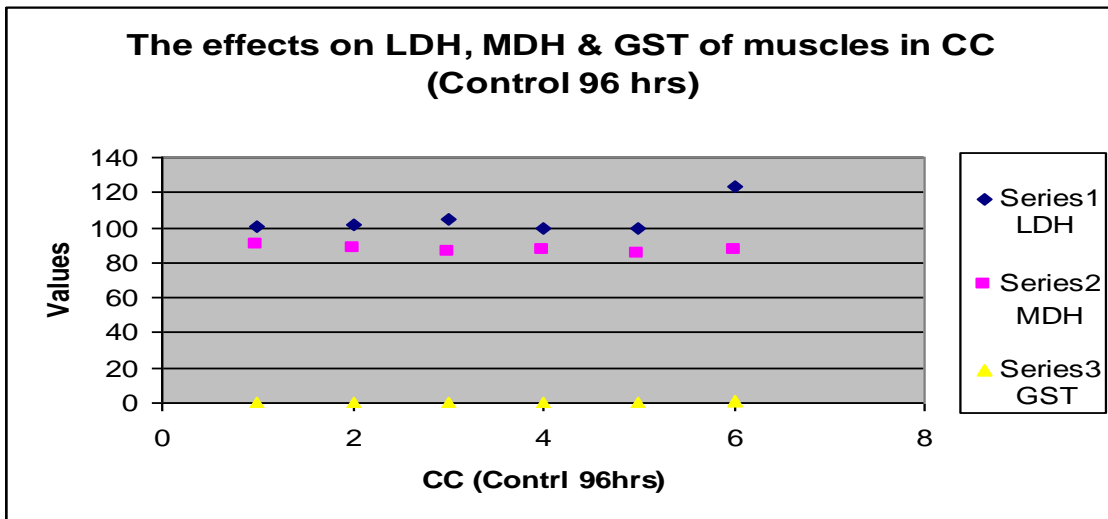


Fig. 5: The effects on LDH, MDH & GST of Skeletal muscles in CC(Control 96hr) of Tambaqui fish.

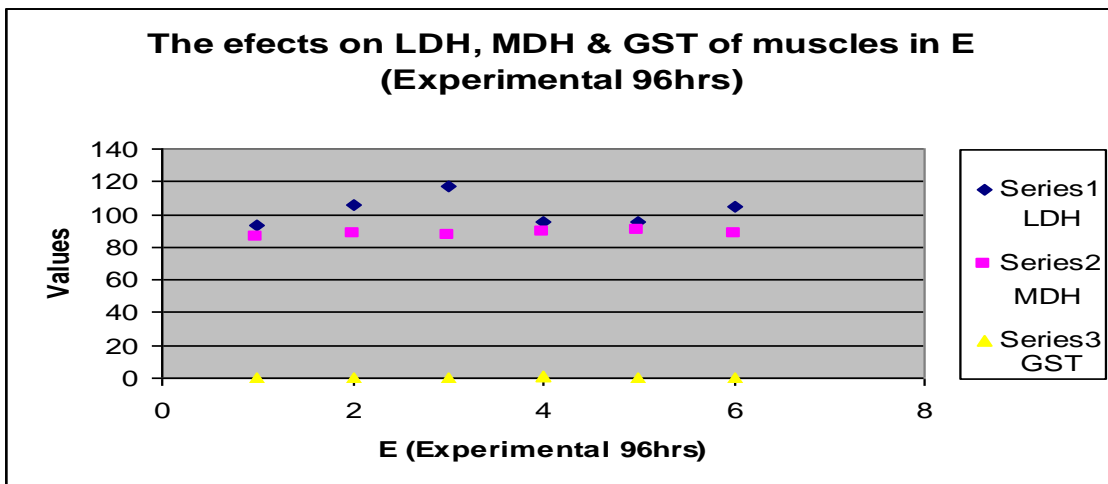


Fig. 6: The effects on LDH, MDH & GST of Skeletal muscles in E(Experimental 96hrs) of Tambaqui fish.

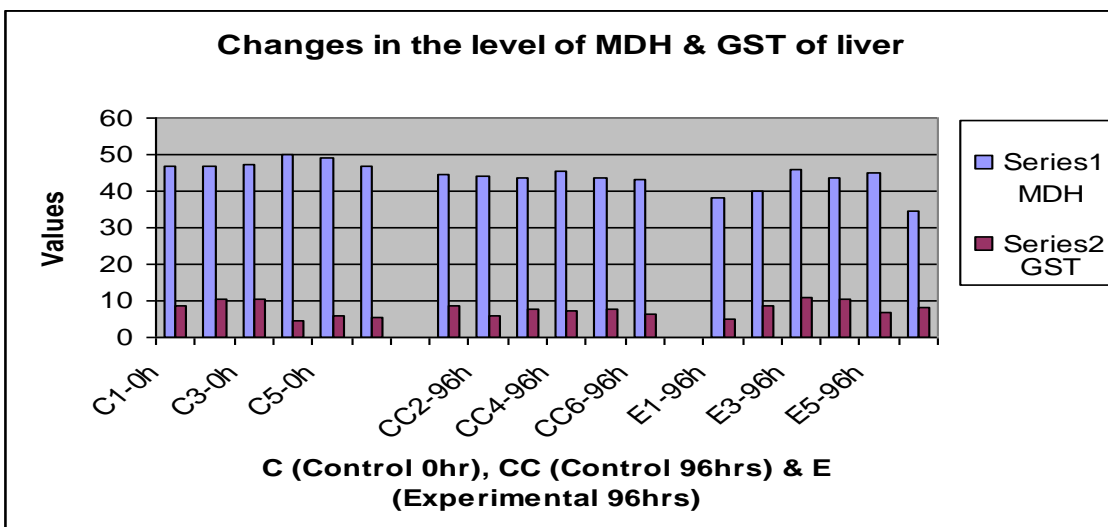


Fig. 7: The effects on MDH & GST of Liver in C(Control 0hr), CC(control 96hrs) and E(Experimental 96hrs) of Tambaqui fish.

Table 2: Change in the level of MDH, LDH & GST (Muscles & Liver) of tambaqui (*Colossoma macropomum*) exposed to different temperature and CO₂.

	LDH Muscle	MDH Muscle	MDH Liver	GST Liver	GST Muscle
C1-0h	100.8334033	88.38434084	46.63384244	8.7571288	0.298881949
C2-0h	102.4762862	90.20799839	46.84485531	10.320564	0.421265604
C3-0h	108.4617926	88.81784566	47.20357717	10.370187	0.321528375
C4-0h	105.6525362	90.19314309	50.0522508	4.3390507	0.470246574
C5-0h	104.1009164	87.48424437	49.06049035	5.854989	0.334895506
C6-0h	113.7823553	92.26607717	46.59163987	5.4118852	0.369363601
Average	105.8845483	89.55894159	47.73110932	7.508967	0.369363601
Std. Dev.	4.677243467	1.696192676	1.464313716	2.6393684	0.065291828
CC1-96h	100.3914791	90.1278135	44.53778135	8.7448199	0.191517023
CC2-96h	101.872916	88.36446342	44.18609325	5.793088	0.208099824
CC3-96h	105.1588505	86.39474277	43.46864952	7.8932844	0.295753861
CC4-96h	99.42926045	86.67986334	45.55767685	7.1798188	0.138006748
CC5-96h	99.38390675	85.55102894	43.67966238	7.529292	0.473998853
CC6-96h	123.244172	87.32134244	43.17323151	6.3460231	0.665035302
Average	104.9134308	87.4065424	44.10051581	7.247721	0.328735268
Std. Dev.	9.234942802	1.632477829	0.866956947	1.0649851	0.202507343
E1-96h	93.25502412	86.51949357	38.36213826	5.0098025	0.470473018
E2-96h	106.2529944	88.59742162	40.02913987	8.6427832	0.412120001
E3-96h	116.7984043	87.39907958	45.68428457	11.014187	0.403209655
E4-96h	95.06846865	88.73057074	43.42644695	10.387174	0.715626912
E5-96h	94.89906752	90.49818328	44.96684084	6.9784706	0.500
E6-96h	104.8638545	87.74566821	34.73271704	8.406	0.500
Average	101.8563023	88.24840283	41.20026125	8.406	0.500
Std. Dev.	9.16626939	1.369600774	4.254529		

3 Results & discussion

The physiological responses, known as stress, are grouped as primary, secondary and tertiary responses (Barton [15]). Primary responses are mediated by the neuron-endocrine system with a fast release of stress hormones; catecholamines (adrenaline and noradrenaline) that are released and synthesized through cromafin cells and cortisol, synthesized by the inter-renals cells. Secondary responses are mediated by the stress hormones and involve the various biochemical and physiological effects associated to stress, which includes metabolic, hematological, hydromineral and structural changes that suggest an important change that helps to evaluate the fish health conditions. Tertiary responses affect the animal as a whole, compromising growth, disease resistance and reproductive success. Fish hematological parameters have been widely used as indicator for stress caused by environmental conditions (Affonso et al. [16], Tavares-Dias & Moraes [17], Carvalho & Fernandes [18]). In order to overcome stress, hemo-concentration is among the important strategies for increasing blood oxygen carrying capacity under high energy demand situations (carvalho & Fernandes [18], Trezado et al. [19], Ziskowski et al. [20]). However, stressors can compromise iron absorption, cause erythrocyte malformation or hemolysis, inhibit hemoglobin synthesis, compete for the oxygen bonding site, resulting in hemodilution or anaemia, reducing the blood oxygen carrying capacity (Affonso [21], Tavares-Dias & Moraes [17]). Therefore, the analysis of hematological parameters in fish can contribute to the assessment of animal health.

The hematological parameters: Ht, Hb, RBC, VCM, HCM & CHCM of the fish from the three treatments are represented in Table 1. The similarity of the Tambaqui physiological profile on all three treatments reflects a satisfactory rearing environment. Similar productive performances in pond systems were obtained by Kubitza et al [22] with *Micropterus salmoides* and Affonso et al., [9] with Tambaqui in pond culture and Good et al. [23] with *O. mykiss*.

In the present study when the Tambaqui fish, exposed to a temperature rise from 28° to 32°C and a CO₂ concentration 10 times higher than today. However we observed a significant increase in (Hb), (HT), (RBC), (VCM), (HCM) & (CHCM) levels in blood, while a significant decrease was observed in Glucose level in blood as compared to control values (Tab. 1) and (Fig. 1, 2 & 3).

Liver MDH values of were 47.731, 44.100 & 41.200 respectively while the values of GST of liver were 7.508 & 7.247 and 8.406 respectively, as compared to control values (Tab. 2). The MDH values in muscles were 89.558, 87.406, 88.248, while the LDH values of muscles were 105.884, 104.913, 101.856, while the GST Values of Muscles were 0.369 & 0.328 respectively as compared to control values (Tab. 2) and (Fig. 4, 5, 6 & 7). Temperature fluctuations are much underappreciated stressor of fish. Similar finding that the slight change in water temperature

and CO₂ can be due to natural heat loading system or thermal pollution in normal or natural water disrupt aquatic ecosystem resulting stress to fish [14].

This study will contribute valuable knowledge needed for fisheries management and aquaculture of Tambaqui fish (*Collosoma macropomum*, Cuvier, 1818) by increasing the knowledge of ecology and environmental effects on the biology of Tambaqui fish (*Collosoma macropomum*).

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