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The cave environment influencing the lipid profile and hepatic lipogenesis of the fish *Ancistrus cryptophthalmus* Reis, 1987 (Siluriformes: Loricariidae)

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Abstract: The metabolism of hypogean organisms is frequently molded by the cave environment traits, especially food scarcity. The aim of the present work was to evaluate the influence of such environment on lipid composition and hepatic lipogenesis in the fish *Ancistrus cryptophthalmus*. For this, the species was compared to an epigeal population of the species. A greater accumulation of total lipids was observed in the cave-dwelling fish (18.36 g/100 g tissue) compared to the surface fish (14.09 g/100 g tissue). The muscle fatty acid profile also varied between the populations. Arachidonic acid was only detected in the epigeal fish, while docosahexaenoic acid was present in the cave fish. In the lipid profile of *Ancistrus cryptophthalmus* there was a higher proportion of saturated fatty acids, followed by monounsaturated and polyunsaturated fatty acids; *Ancistrus* sp. showed a predominance of monounsaturated fatty acids. Significant differences were also observed in the activities of the hepatic enzymes glucose-6-phosphate dehydrogenase and malic enzyme. The activities of these two enzymes were greater in the epigeal animals. The differences could be related to different food availability observed in the two environments. An ecotone zone was observed, located next to the entrance of the Lapa do Angélica cave (Goiás State, Brazil), where the fishes showed characteristics that were intermediate between those of hypogean fishes from deeper within the cave, and the epigeal population. It could be concluded that the characteristics of the cave environment significantly influenced the composition of muscle fatty acids and lipogenesis in the hypogean fish *Ancistrus cryptophthalmus*.

Keywords: fish; cave; fatty acids; malic enzyme; glucose-6-phosphate dehydrogenase; metabolism

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INTRODUCTION

Fish are important components of subterranean aquatic communities, where they often are at the top of the food web. Globally, the number of known fish species is currently around 28,000, of which 299 can be found in the hypogean environment, with at least 164 showing some degree of troglomorphism (Romero & Paulson, 2001; Romero, 2009; Proudlove, 2010).

Brazil possesses a rich subterranean ichthyofauna, with more than 25 fish species that present troglomorphism compared to the corresponding epigeal species (Trajano & Bichuette, 2010). Most of these troglobitic fishes belong to the Siluriformes, especially the Trychomictoridae family, although there are representatives of other families and also of other orders (such as the Characiformes and Gymnotiformes)

(Trajano & Bichuette, 2010). Globally, there are some troglobitic armored catfishes that belong to the Loricariidae family (*Ancistrus*). There are two Brazilian fishes of this genus: *Ancistrus cryptophthalmus*, found in the State of Goiás, central Brazil (Sabino & Trajano, 1997), and *Ancistrus formoso*, found in the State of Mato Grosso do Sul (Trajano & Bichuette, 2010).

Although the diversity of hypogean fishes is fairly well known, there exists little information concerning the metabolism of these organisms. Nevertheless, it is believed that the animals are hypometabolic, and tend to develop elaborate mechanisms for both searching for food and improving the capacity for lipid storage (Wilkens, 2005; Salin et al., 2010).

Lipids are important components of the diets of many animals, and are an efficient source of both energy and the fatty acids that are essential for

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development. In ectothermic animals, such as fish, lipids act as structural components responsible for the maintenance of fluidity of the plasma membrane, as well as for the storage and supply of energy, and are therefore essential for basic physiological functions (Brown & Murphy, 1995). An ability to synthesize a wide variety of fatty acids is essential for all living organisms.

In fish, biosynthesis of fatty acids occurs mainly in the liver, and proceeds by the successive addition of two carbon units, derived from the malonyl group, to the growing lipid chain (Nelson & Cox, 2010). This process involves the action of specific enzymes such as glucose-6-phosphate dehydrogenase, which provides energy in the form of NADPH in the pentose phosphate pathway, and the malic enzyme, which is involved in the pyruvate/malate cycle (Wang et al., 2005).

In most fish species, the fatty acid profile and the activity of the enzymes concerned with lipogenesis are affected by the composition of the diet and the frequency of feeding (Visentainer et al., 2005; Wang et al., 2005; Visentainer et al., 2007). For this reason, subterranean environments, which tend to be strongly oligotrophic, can promote alterations such as diminution of metabolism in cave fish, and a tendency for greater storage of lipids (Wilkins, 2005).

The objective of this work was therefore to investigate the influence of the subterranean environment on the lipid profile and hepatic lipogenesis of the fish *Ancistrus cryptophthalmus*. This was achieved using comparisons with epigeal fish that inhabited the external stretch of the same water channel present in the Lapa do Angélica cave (Goiás State, Brazil). In addition, the study sought to identify whether there were any differences between individuals of the hypogean fishes that inhabited zones closer to the cave entrance (where there was a greater supply of nutrients), and fish that were found deeper within the cave.

Study region

The São Domingos karst region is located in the Serra do Calcário, in the central-west Brazilian State of Goiás, where the predominant vegetation is that of the cerrado (Brazilian savannah). This area has a semi-humid tropical climate, with a dry period that extends from April to September (Klink & Machado, 2005). The region is known for its rich subterranean ichthyofauna, and 66 limestone caves have been recorded, amongst which is the Angélica cave in the Terra Ronca State Park (13° 31' 22.2" S, 46° 22' 55.5" W).

The Angélica cave is the fourth largest Brazilian cave in terms of extent, with 14,100 meters of mapped galleries, 8,000 meters of which contain large, fast-flowing watercourses (Auler et al., 2001). The stream conduit has an average width of around 5 meters, and the water depth varies between 0.5 and 2 meters. In the dry season, the measured discharge of water at the mouth of the cave was 2.27 m³/s (Trajano & Bichuette, 2007).

Fish species studied

The *Ancistrus cryptophthalmus* Reis, 1987 (Fig. 1A) is found in four caves in the São Domingos karst region

(the Angélica, Bezerra, São Vicente, and Passa Três caves). They have poorly-developed ocular structures, reduced body pigmentation, detritivore feeding habits, and precocial lifestyles. These characteristics can be interpreted as being specialized adaptations to the subterranean environment (Secutti & Trajano, 2009).

In addition to the hypogean populations, the same region harbors epigeal populations of the genus *Ancistrus*, which include an unidentified species (Fig. 1B). These animals, which are morphologically distinct to the hypogean, have better developed ocular structures, and intense pigmentation (Reis et al., 2006).

The fish of the genus *Ancistrus* provide an excellent model for studies concerning the adaptive evolution of cave-dwelling groups, because there are relatively abundant populations of the animals inhabiting both hypogean and epigeal environments in the same region (Trajano & Bichuette, 2007).

METHODS

Capture and transport

Ten individuals of *Ancistrus cryptophthalmus* (hypogean) and *Ancistrus* sp. (epigeal) were collected manually using a net, in May 2011. No consideration was given to differences between the animals in terms of physiological factors, age, size, weight, stage of maturity, or anatomical features, in order to obtain a sample that was more representative of the overall population. In addition, the fish were handled as little as possible to minimize situations of stress.

The hypogean fish were collected at two points within the Angélica cave. Five individuals were obtained near the entrance (site P, located around 300 meter far from the entrance), and the remainder deeper within the cave, approximately 1000 meters from the entrance (site P2) (Fig. 2). Since fish of the species *Ancistrus cryptophthalmus* are highly sedentary and live their entire life in a stretch of water less than 100 m long, there was a low probability of movement of the animals between the two locations (Trajano & Bichuette, 2007). The non-troglobitic fish were collected in the epigeal stretch

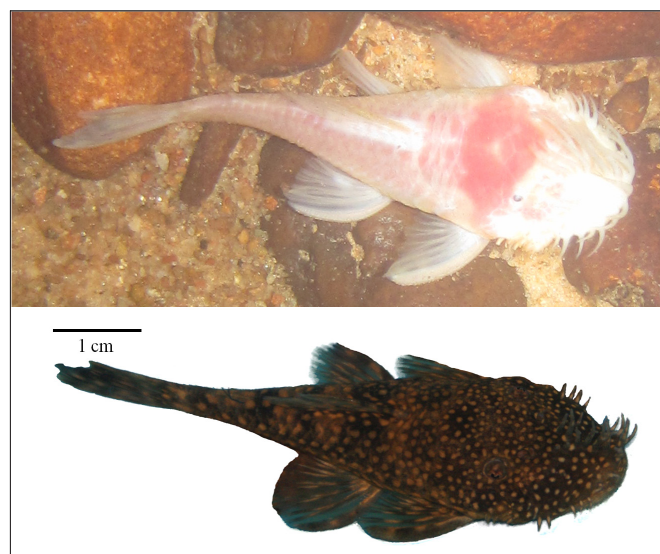


Fig. 1. Species of fish studied: A) *Ancistrus cryptophthalmus*; B) *Ancistrus* sp.

of the Angélica River, at approximately 300 meters from the cave entrance.

Measurements of water pH, temperature, and dissolved oxygen content were made *in situ* at the time of collection, using a multiprobe sensor (YSI). Samples of substrate were also collected from the riverbed at the three collection points, for later analysis of organic matter using the procedure described by Davies (1974).

The animals were transported to Universidade Federal de Lavras, in groups of five, in plastic containers (5 liter capacity) protected from light and with constant aeration. The containers were filled to the rim with water from the Angélica river in order to avoid impacts that could cause a situation of stress to the fish, according to the norms established by the Commission for Ethics in the Use of Animals of the Federal University of Lavras (license number: 002/11).

Genetic analysis

Samples of fins and muscle were obtained from 27 fishes representing four populations (22 from Angélica Cave: 5 specimens from a hypogean population inside the cave, 5 from a hypogean population near the surface and 11 from an epigeon population outside the cave and described only as *Ancistrus* sp.). Five hypogean individuals from Passa-Três Cave were also analyzed. All tissues were stored in 95% ethanol. DNA was isolated

and 658 base pairs of the mitochondrial gene *Cytochrome c Oxidase I* (COI) amplified using the primers FishF1 and FishR1 (Ward et al., 2005) as described in Carvalho et al. (2011). Sequences were determined bi-directionally using the BigDye Terminator v3.1 Cycle Sequencing Kit (Applied Biosystems, Inc., Foster City, California, USA) following the manufacture's protocol on an Applied Biosystems Inc. 3130 Genetic Analyzer. Each recovered haplotype was deposited in GenBank (Accession numbers: JX477622-JX477648). One previously described *Ancistrus brevipinnis* COI sequence (Genbank number: EU359402.1) was also included in the analysis.

DNA sequences were analyzed using SeqScape v2.7[®] software (Applied Biosystems, Inc., Foster City, California, USA). Genetic distances were estimated using Kimura's two-parameter (K2P) nucleotide substitution model (Kimura, 1980) using MEGA version 5.10 (Tamura et al., 2011).

Biochemical analysis

After transport, the animals were desensitized in an ice bath, weighed, and eviscerated, with separation of the liver for use in the enzymatic analyses.

Samples of muscle tissue were submitted to lipid extraction and esterification using the method of Folch et al. (1957). The methyl esters of the fatty acids were determined using a Shimadzu GC-2010

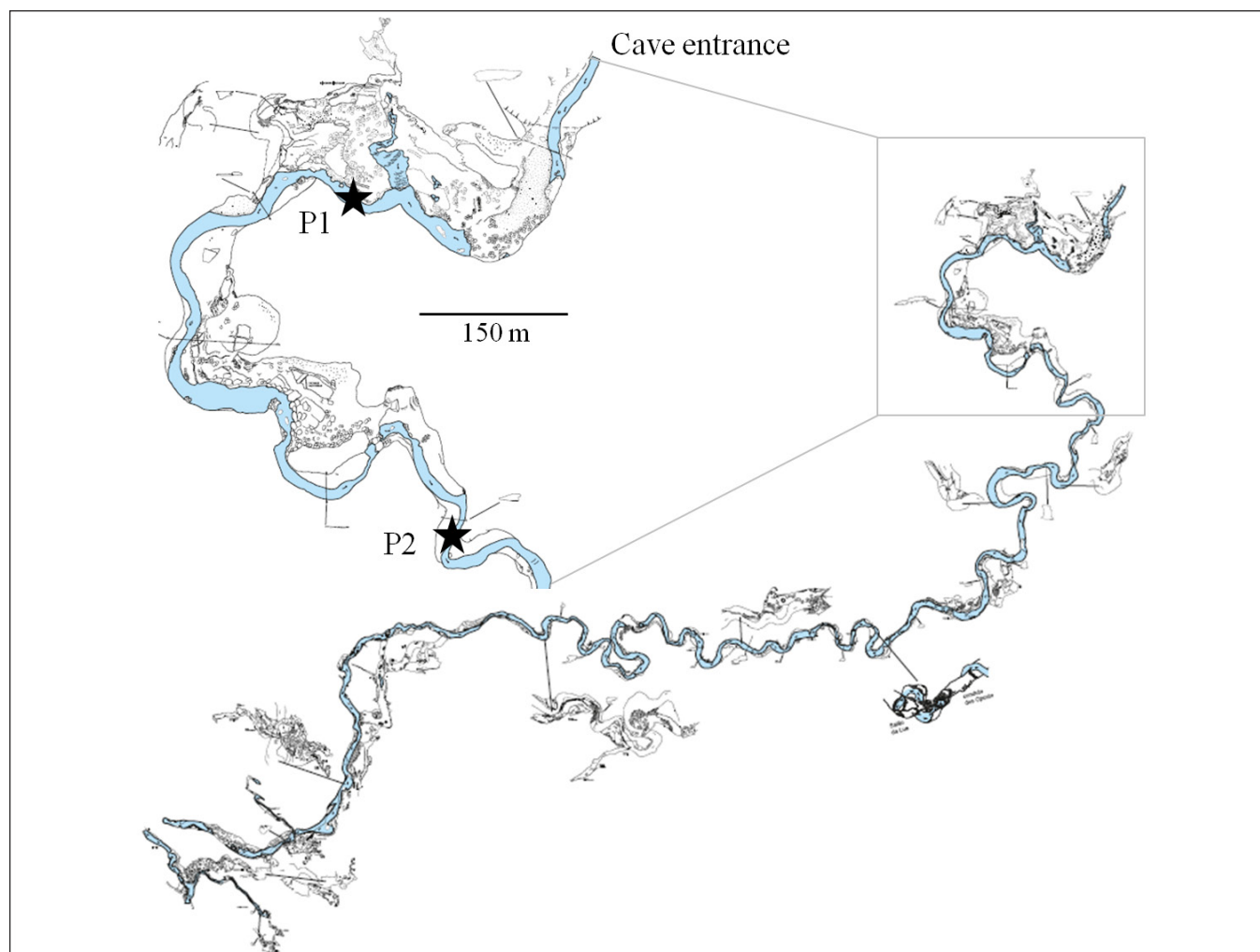


Fig. 2. Map of the Angélica Cave, indicating collection points P1 and P2.

gas chromatograph equipped with a flame ionization detector and a DB-WAX fused silica capillary column (100 m × 0.25 mm). The operating conditions were as follows: splitless injection mode; 1 µL injection volume; 260 °C detector temperature; 260 °C injector temperature; temperature program: 4 °C min⁻¹ to 140 °C, hold at 140 °C for 5 minutes, ramp at 4 °C min⁻¹ to 240 °C, hold at 240 °C for 30 minutes.

The chromatographic peaks were identified by comparing retention times with those of a commercial standard (Supelco 37 Component FAME Mix, supplied by Sigma-Aldrich). The fatty acids were quantified by normalization of the peak areas, and the values were converted to percentages of the mass of muscle tissue, multiplying by the total lipid fraction content and by the conversion factor for fatty fish (0.9), as described by Holland et al. (1994).

The samples of liver tissue were homogenized in 25 mM HEPES-KOH buffer at pH 7.4, centrifuged at 10,000 rpm (4 °C for 30 minutes), and analyzed for the specific enzymatic activities of glucose-6-phosphate dehydrogenase and malic enzyme, according to the methods of Graeve (1994) and Spina et al. (1966), respectively. The enzyme kinetics was monitored using UV spectrophotometry (Shimadzu UV-1800). The total soluble protein contents of the livers were determined using the method described by Bradford (1976).

All the results were tested statistically using analysis of variance (ANOVA) and Tukey's test to identify significant differences among the sample mean values. The software used was R (R Development Core Team, 2011), and the significance level employed was 5%.

RESULTS

At the time of sample collection, the river water at the epigeal site presented a temperature of 24.8 °C, a dissolved oxygen content of 94.7%, and a pH of 6.3. Inside the cave, where there was no significant difference between the conditions at the two collection points, the temperature was higher (25.3 °C), and the dissolved oxygen content was lower (91.5%). The water pH was the same at the hypogean and epigeal sites. Physico-chemical information concerning the collection points is important because the conditions can significantly influence the metabolism of the fish.

The organic matter content of the sedimented substrate differed substantially between the two environments, with values of 1.84 and 0.38 g/cm² obtained at the epigeal site and hypogean site P1 (closer to the cave entrance), respectively. The organic matter content was below the detection limit at the second hypogean site (deeper within the cave).

After trimming unclear ends, a total of 556 base pairs (bp) were obtained from each specimen. Three haplotypes were recovered from all samples analyzed, with a K2P genetic divergence ranging from 0 to 0.2%. When comparing fishes collected only at Angélica Cave (5 individuals from a hypogean population inside the cave, 5 from a hypogean population near the surface and 11 from an epigeal population) only one haplotype was recovered and no genetic divergence was observed between populations. At Passa-Três

Cave, two polymorphic sites were recovered in two specimens, resulting in a divergence of 0.2% between haplotypes. When comparing *Ancistrus* sp and *Ancistrus cryptophthalmus* with *Ancistrus brevipinnis* a divergence of 6.9-7.2% was recovered.

The *Ancistrus cryptophthalmus* fish presented higher body weights, with values of between 2.09 and 2.17 g, and greater deposition of lipids in the muscle tissue (17.6-18.36%), compared to the epigeal population whose average weight and muscle total lipid content were 1.21 g and 14.09%, respectively (Fig. 3).

The lipid profiles revealed the presence of 14 fatty acids in the *Ancistrus cryptophthalmus* individuals collected at site P1, and 10 in fish of the same species collected at site P2. The lipid profile of the epigeal *Ancistrus* sp. revealed the presence of 19 fatty acids (Table 1). Amongst those fatty acids identified in both species, the most prevalent were myristoleic (C14:1), palmitic (C16:0), palmitoleic (C16:1), oleic (C18:1ω9c), and linoleic (C18:2ω6c) acids.

Significant differences between the hypogean and epigeal fishes were observed in the proportions of saturated and monounsaturated fatty acids. While the cave fish from both sites within the cave showed high levels of saturated fatty acids (62.88% for P1, and 62.75% for P2), and smaller proportions of monounsaturated fatty acids (24.54% for P1, and 24.76% for P2), the epigeal fish showed a higher content of monounsaturated (46.86%) than saturated (39.22%) fatty acids.

The total content of polyunsaturated fatty acids showed no significant difference between the populations, with the omega-6 series predominant over the omega-3 series in both cases.

In the analysis of the rates of lipogenesis, it was found that the specific activities of the hepatic enzymes (glucose-6-phosphate dehydrogenase and malic enzyme) differed significantly ($P < 0.05$) between the hypogean and epigeal fishes (Table 2). Higher specific enzyme activities in the epigeal fish were indicative of greater lipogenesis in this population.

DISCUSSION

Studies of metabolism are essential for understanding the basic needs of animals, and were used here to elucidate the habits of the

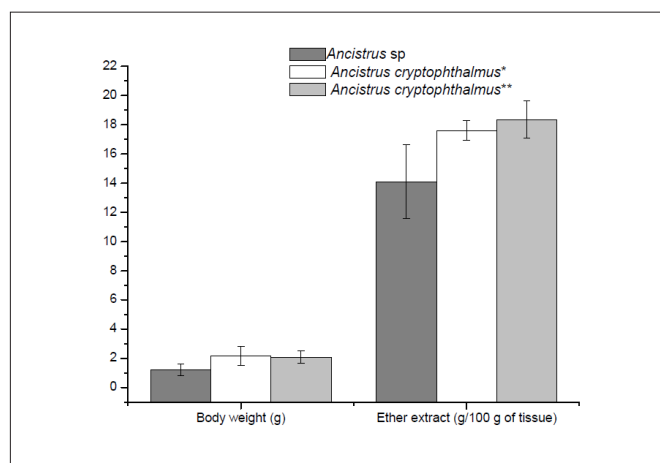


Fig. 3. Body weights and ether extract amounts of the epigeal and troglitic fish.

* Collected at site P1.

** Collected at site P2.

Table 1. Fatty acid profiles of the species *Ancistrus cryptophthalmus* and *Ancistrus* sp.

Values are shown as means \pm standard deviation. Where the values expressed in % (on the same line) are followed by identical letters, differences between the values were not significant ($p > 0.05$). Where the values expressed in g/100 g (on the same line) are followed by identical Greek letters, differences between the values were not significant ($p > 0.05$). Differences were evaluated using analysis of variance (ANOVA) and Tukey's test. nd = not determined.

Fatty acid	<i>Ancistrus</i> sp.		<i>Ancistrus</i> cryp.*		<i>Ancistrus</i> cryp.**	
	%	g/100g	%	g/100g	%	g/100g
C14:0	0.81 \pm 0.10b	0.10 \pm 0.01 β	5.01 \pm 0.61a	0.85 \pm 0.10 α	5.93 \pm 1.59a	0.96 \pm 0.25 α
C14:1	13.78 \pm 1.73a	1.79 \pm 0.22 α	nd	nd	14.22 \pm 3.06a	2.27 \pm 0.50 α
C15:0	0.80 \pm 0.08c	0.10 \pm 0.01 γ	4.58 \pm 0.70a	0.77 \pm 0.11 α	3.35 \pm 0.25b	0.53 \pm 0.04b
C 15:1	0.94 \pm 0.15b	0.12 \pm 0.01 β	4.89 \pm 0.56a	0.82 \pm 0.09 α	nd	nd
C16:0	25.67 \pm 1.66b	3.30 \pm 0.21 β	48.82 \pm 1.01a	8.18 \pm 0.19 α	49.54 \pm 5.69a	7.78 \pm 0.88 α
C16:1	11.36 \pm 0.40a	1.45 \pm 0.05 α	1.88 \pm 0.91b	0.31 \pm 0.15 β	10.54 \pm 2.60a	1.64 \pm 0.40 α
C17:0	0.46 \pm 0.10c	0.06 \pm 0.01 γ	3.50 \pm 0.61a	0.58 \pm 0.10 α	1.55 \pm 0.35b	0.24 \pm 0.05 β
C17:1	0.81 \pm 0.15b	0.10 \pm 0.01 β	2.41 \pm 0.45a	0.39 \pm 0.07 α	nd	nd
C18:0	8.06 \pm 0.52	1.01 \pm 0.06	nd	nd	nd	nd
C18:1 ω 9c	19.40 \pm 0.56a	2.43 \pm 0.07 α	11.42 \pm 0.65b	1.86 \pm 0.10 β	nd	nd
C18:2 ω 6t	1.69 \pm 0.34	0.21 \pm 0.04	nd	nd	nd	nd
C18:2 ω 6c	7.07 \pm 0.60a	0.88 \pm 0.07 α	5.61 \pm 1.155 a	0.91 \pm 0.18 α	7.01 \pm 1.48a	1.06 \pm 0.22 α
C20:0	2.77 \pm 0.21a	0.34 \pm 0.02 α	0.95 \pm 0.24b	0.15 \pm 0.03 β	2.35 \pm 0.31a	0.35 \pm 0.04 α
C18:3 ω 6	0.51 \pm 0.36b	0.19 \pm 0.04 α	0.95 \pm 0.41a	0.21 \pm 0.06 α	1.09 \pm 0.34a	0.16 \pm 0.05 α
C20:1	0.51 \pm 0.004	0.06 \pm 0.004	nd	nd	nd	nd
C22:0	0.63 \pm 0.13	0.07 \pm 0.01	nd	nd	nd	nd
C20:3 ω 6	1.98 \pm 0.32	0.24 \pm 0.03	nd	nd	nd	nd
C20:4 ω 6	0.38 \pm 0.007	0.04 \pm 0.008	nd	nd	nd	nd
C22:2	nd	nd	0.78 \pm 0.11	0.13 \pm 0.09	nd	nd
C24:1	nd	nd	3.92 \pm 0.75	0.56 \pm 0.10	nd	nd
C22:6 ω 3	1.18 \pm 0.20b	0.14 \pm 0.02 β	4.85 \pm 0.66a	0.74 \pm 0.10 α	4.36 \pm 0.95a	0.63 \pm 0.13 α
Σ Saturates	39.22 \pm 1.98b	5.00 \pm 0.25 β	62.88 \pm 1.07a	10.55 \pm 0.18 α	62.75 \pm 6.54a	9.86 \pm 1.02 α
Σ Monounsaturates	46.83 \pm 1.47a	5.96 \pm 0.19 α	24.54 \pm 1.24b	3.96 \pm 0.20 β	24.76 \pm 5.06b	3.90 \pm 0.81 β
Σ Polyunsaturates	13.94 \pm 1.10a	1.71 \pm 0.13 α	12.56 \pm 0.48a	2.00 \pm 0.07 α	12.47 \pm 2.14a	1.86 \pm 0.31 α
Σ ω 3	1.18 \pm 0.20b	0.14 \pm 0.02 β	4.85 \pm 0.66a	0.74 \pm 0.10 α	4.36 \pm 0.95a	0.63 \pm 0.13 α
Σ ω 6	12.75 \pm 1.14a	1.57 \pm 0.14 α	6.94 \pm 1.12b	1.12 \pm 0.18 β	8.10 \pm 1.80b	1.22 \pm 0.26 β

* Collected at site P1.

** Collected at site P2.

cave fish. The hypogean environment is unusual when compared to epigeal environments, since there is no photoperiod and food availability is low. For many animals, these factors preclude the establishment of viable populations in subterranean ecosystems (Culver et al., 2009). A fish species would become actually established in the hypogean environment, if it develop morphological, physiological, metabolic, and behavioral specializations (Mathieu & Hervant, 2006; Reis et al., 2006).

In a study of the epigeal fish species *Micropterus salmoides*, it was shown that when subjected to periods of fasting, the animals developed the capacity to store proteins and lipids as a means of guaranteeing a supply of energy (Cyrino et al., 2000). This ability to store nutrients, developed by various fish species as a result of fasting (Salin et

al., 2010), provides a basis for an understanding of the results obtained in the present work. The hypogean fishes presented a higher weight and greater lipid deposition in the muscle tissue than the epigeal *Ancistrus* sp., which is provided with a greater quantity of organic matter in its habitat.

A large sinkhole at the entrance to the Angélica cave enables transport of quantities of organic matter, mainly leaves and small invertebrates, into the cave interior. However, fish of the *Ancistrus* genus are essentially grazers, so that even given the transfer of such organic matter, the amount of food available to these fish is considerably smaller in the cave habitat, and consists only of a film of fine particulate organic matter (Power, 1990; Trajano & Bichuette, 2007).

It is recognized that a reduction in the availability of food is generally associated with a worsening of the

Table 2. Specific activities of glucose-6-phosphate dehydrogenase and the malic enzyme in epigeal and hypogeal fish. Values are shown as means \pm standard deviation. Where the values on the same line are followed by identical letters, differences between the values were not significant ($p > 0.05$). Differences were evaluated using analysis of variance (ANOVA) and Tukey's test.

Enzyme	<i>Ancistrus</i> sp. (U/mg of ptn)	<i>Ancistrus cryp.*</i> (U/mg of ptn)	<i>Ancistrus cryp.**</i> (U/mg of ptn)
Glucose-6-phosphate dehydrogenase	0.633 \pm 0.116a	0.271 \pm 0.097b	0.189 \pm 0.098b
Malic	0.048 \pm 0.011a	0.034 \pm 0.006a	0.013 \pm 0.011b

* Collected at site P1.

** Collected at site P2.

condition of fish (Oscoz et al., 2005). Nonetheless, metabolic changes, such as greater deposition of body fat, occur in order to maintain a constant nutritional status and bodily condition, even in an environment with limited resources, such as a cave.

Although the total lipid contents of the muscle tissues of the *Ancistrus cryptophthalmus* fish were higher, the variety of fatty acids in the lipid profile was smaller than observed in the epigeal population. Several studies have shown that the profile of fatty acids found in the muscle tissues of fish reflects the variability of these compounds in the diets of the animals (Visentainer et al., 2005; Vieira et al., 2011). Valente et al. (2007) found that in the fish species *Dicentrarchus labrax*, the provision of linoleic acid in the diet was directly reflected in the muscle content of this fatty acid as well as its derivatives formed during the processes of elongation and desaturation. The findings of the earlier studies were corroborated by the results obtained in the present work, where the hypogeal fish that had a more restricted diet (in an habitat poor in organic matter) exhibited a reduced range of fatty acids in their muscle tissues, compared to the epigeal organisms.

Regions close to the entrances of caves often show structural, biological, and physical gradients, creating a zone of transition between the epigeal and hypogeal systems. Such regions can be considered as ecotones, with distinct characteristics due to the equilibrium established between the epigeal zone, where resources are readily available, and the hypogeal zone, where the environment is more stable (Prous et al., 2004).

Due to the existence of an ecotone zone located close to site P1, the hypogeal fishes collected at this area presented a range of fatty acids that was intermediate between those of the epigeal fish and the animals collected at the site deeper within the cave. It is therefore possible that the diet of the fish from site P1 was richer and more diversified than that of fish from deeper regions of the cave, but less rich than that of their epigeal population. Even though the sample population was small (5 fish at each collection point), the information presented is representative of the total population of *Ancistrus cryptophthalmus* in the Angélica cave, since it employed data that were biochemical in nature, rather than morphological measurements alone.

The elevated levels of palmitic (C16:0) and palmitoleic (C16:1) acids in both of the population studied, and the lower levels of long chain fatty acids (C20 and C22), are common characteristics of freshwater fish (Çelik et al., 2005; Inhamuns et al., 2009). On the other hand, stearic acid (C18:0), which is also common in the lipid profile of freshwater fish, was not found in the hypogeal fish *Ancistrus cryptophthalmus*. Since the fatty acid composition reflects the diet of the animal, it can therefore be inferred that the diet of the cave-dwelling fish was probably deficient in this fatty acid (Inhamuns et al., 2009).

Eicosenoic (C20:1), behenic (C22:0), eicosatrienoic (C20:3 ω 3), and arachidonic (C20:4 ω 6) acids were only detected in the epigeal fish. Arachidonic acid (C20:4 ω 6), which is nutritionally and physiologically important for fish, is synthesized from the essential linoleic acid (C18:2 ω 6) present in abundance in cave-dwelling fish (Martino, 2003). The absence of arachidonic acid in individuals of the *Ancistrus cryptophthalmus* is suggestive of a deficiency in the specific desaturases and elongases responsible for catalyzing the conversion of linoleic acid into longer chain derivatives (such as arachidonic acid).

It should also be noted that conditions of stress, of any nature, predispose fish to various illnesses. Hence, in an environment where conditions are adverse, such as that of a cave, compared to epigeal environments a fraction of the polyunsaturated fatty acids that are synthesized is directed to the immunological system in order to maintain it active, which could explain the low availability of these compounds in the muscle tissues of *Ancistrus cryptophthalmus* (Sargent et al., 1999).

In fish, the balance between saturated and unsaturated fatty acids is intimately related to maintenance of the fluidity of the cell membranes, and is greatly influenced by temperature (Ribeiro et al., 2011). Generally, a decrease in temperature results in an increase in the degree of unsaturation, since unsaturates are required in the membrane phospholipids to maintain flexibility and permeability (Lovell, 1991). Conversely, in higher temperature aquatic environments, it is often the saturated fatty acids that predominate in the lipid profiles (Visentainer et al., 2007).

This is supported by the present findings, where *Ancistrus cryptophthalmus* showed higher levels of saturated fatty acids, while monounsaturates predominated in the epigeal fishes, since during the collection period, which took place on a sunny day in the dry season, the hypogeal site on the Angélica river was 0.5 °C warmer than the epigeal site.

Meanwhile, it is important to note that according to Secutti & Trajano (2009) there is a diurnal temperature cycle in the Angélica cave, within a range of around 22.5 °C to 25 °C, and that seasonal changes in climate can affect the lipid profile of fish (Inhamuns et al., 2009; Kalyoncu et al., 2009). Further work over a more extended period will be needed to fully investigate these effects.

As observed for *Ancistrus cryptophthalmus*, other studies of freshwater fish have also shown that when

the proportion of saturated fatty acids in the muscle tissues increases, the level of polyunsaturated fatty acids decreases (Inhamuns et al., 2009).

In addition to *Ancistrus* sp., other fish that are not cave-dwellers, such as cachara (*Pseudoplatystoma fasciatum*), pintado (*Pseudoplatystoma corruscans*), pacu (*Piaractus brachipomus*), and dourado (*Salminus brasiliensis*) present higher concentrations of monounsaturated fatty acids in their muscle tissues (Ramos Filho et al., 2008; Sharma et al., 2010).

Considering the polyunsaturated fatty acids, Gutierrez & Silva (1993) reported that the great majority of freshwater fish caught in Brazil are deficient in the omega-3 series fatty acids, such as eicosapentaenoic acid (C20:5 ω 3) and docosahexaenoic acid (C22:6 ω 3). These two compounds are produced endogenously from α -linolenic acid, found in large quantities in marine plants.

Eicosapentaenoic acid (C20:5 ω 3) was not detected in either *Ancistrus cryptophthalmus* or *Ancistrus* sp., while docosahexaenoic acid (C22:6 ω 3) was found at higher concentrations in the troglobitic fish.

According to Sargent et al. (1999), docosahexaenoic acid (C22:6 ω 3) is extremely important for animal development, and can be found at high levels in neural tissues as well as the ocular apparatus of fish. Since the ocular structure of *Ancistrus cryptophthalmus* is less developed than that of the corresponding epigeal organisms, there is greater availability of this fatty acid in the muscle tissue.

The greater part of lipid synthesis occurs in the hepatic tissue of fish (Figueiredo-Silva et al., 2005; Valente et al., 2007). In most animals, the pentose phosphate pathway, catalyzed by the glucose-6-phosphate dehydrogenase enzyme, is the principle source of NADPH (up to 60%) for the synthesis of fatty acids. The malic enzyme, important in the pyruvate/malate cycle, provides an alternative route that can also contribute to the supply of NADPH. The contribution of each of these enzymes depends, amongst other factors, on the species and the conditions to which the fish are subjected (Wang, 2005).

The specific enzyme activity values obtained here for both fish populations are in agreement with data reported in the literature, according to which specific activity values for fish lie in the ranges 0.109-0.920 U/mg of protein for glucose-6-phosphate dehydrogenase, and 0.013-0.082 U/mg of protein for the malic enzyme (Borba et al., 2003; Valente et al., 2007).

It has been reported that the activities of the enzymes that participate in the processes of lipogenesis and lipid storage in fish (such as glucose-6-phosphate dehydrogenase and the malic enzyme) are affected by the composition of the diet of the animal (Valente et al., 2007). Studies have shown that in many species of fish lipogenesis is stimulated by diets rich in carbohydrates (Dias et al., 2004), and is suppressed when the diet contains a high lipid content (Catacutan & Coloso, 1997; Wang et al., 2005).

The specific activities of the two enzymes investigated were lower in the *Ancistrus cryptophthalmus* collected at site P2, compared to the epigeal population, which could have been due to a scarcity of food in the cave environment (Catacutan & Coloso, 1997; Dias et al., 2004).

In the case of the malic enzyme, the specific activities were the same for *Ancistrus cryptophthalmus* collected at site P1 and the *Ancistrus* sp.. Site P1 was located close to the entrance of the Angélica cave, in an area that could be characterized as a transition zone, and where the concentration of organic matter in the sedimented material was intermediate between the concentrations measured at site P2 and the epigeal site. Hence, the hypogean fish found in this region of the cave retained the principal route for production of NADPH (the pentose phosphate pathway), while at the same time the greater availability of food induced higher activity of the malic enzyme in the alternative route.

The mtDNA COI region could correctly differentiate *Ancistrus brevipinnis* from *Ancistrus* sp and *A. cryptophthalmus*, but could not differentiate the epigeal species *Ancistrus* sp from the hypogean species *A. cryptophthalmus*. The lack of genetic divergence observed between *Ancistrus* sp and *A. cryptophthalmus* is strong evidence that both morphotypes correspond to the same species. Phenotypic plasticity and gene flow between epigeal and hypogean populations are possible hypotheses that should be tested in order to explain the contradiction between morphological and molecular data.

The most remarkable ecological characteristics presented by hypogean fish populations are their small size, restricted geographical distribution, vulnerability to environmental disturbances and sensitivity to stress (Bichuette & Trajano, 2008). Due to the fragility of these populations, it is very important to elucidate the metabolism of cave fishes in order to understand their habits and create favorable conditions for their preservation.

CONCLUSIONS

It is concluded that the characteristics of the cave environment, especially the quantity of available food, had a significant influence on the composition of fatty acids in the muscles, as well as on lipogenesis, in the fish *Ancistrus cryptophthalmus*, which contributed to differences between the hypogean and epigeal populations.

The existence of an ecotone zone was observed close to the entrance of the Angélica cave, where the characteristics of the fish were intermediate between those of hypogean and epigeal populations.

Since *Ancistrus cryptophthalmus* and *Ancistrus* sp. (undescribed and non-troglophobic species) comprise the same species (as shown by the genetic analysis), we cannot consider *A. cryptophthalmus* as a troglobitic species, but as a trogliphilic species with troglomorphic populations.

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