

Trypanosomatids (Protozoa: Kinetoplastida) in three species of Armored Catfish from Mogi-Guaçu river, Pirassununga, São Paulo, Brazil

Tripanossomatídeos (Protozoa: Kinetoplastida) em três espécies de cascudos do rio Mogi-Guaçu, Pirassununga, São Paulo, Brasil

Julia Pereira Molina¹; Rubens Riscala Madi²; Vera Nisaka Solferini³; Paulo Sérgio Ceccarelli⁴; Hildete Prisco Pinheiro⁵; Marlene Tiduko Ueta^{1*}

¹ Departamento de Biologia Animal, Instituto de Biologia, Universidade Estadual de Campinas – UNICAMP, Campinas, SP, Brasil

² Laboratório de Biologia Tropical, Instituto de Tecnologia e Pesquisa, Universidade Tiradentes – UNIT, Aracaju, SE, Brasil

³ Laboratório de Diversidade Genética, Departamento de Genética e Evolução e Bioagentes, Instituto de Biologia, Universidade Estadual de Campinas – UNICAMP, Campinas, SP, Brasil

⁴ Centro de Pesquisa e Gestão dos Recursos Pesqueiros Continentais – CEPTA, Instituto Chico Mendes de Conservação da Biodiversidade – ICMBio, Pirassununga, SP, Brasil

⁵ Instituto de Matemática, Estatística e Computação Científica, Universidade Estadual de Campinas – UNICAMP, Campinas, SP, Brasil

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Abstract

Trypanosome infections have been reported in several species of fish, in majority of cases described on the basis of morphological characteristics. Trypanosomes in fish are heteroxenous and transmitted by hirudineans. This study aims to evaluate the prevalence and density of infections by *Trypanosoma* sp. in blood from three species of catfish, *Hypostomus regani*, *H. strigaticeps*, *H. albopunctatus*, from the Mogi Guaçu River, Pirassununga, São Paulo, Brazil. Further, this study intends to characterize the *Trypanosoma* specimens found in the blood of these fish by morphological and molecular techniques. The trypanosomes overall prevalence observed was 47.6% with a general average density of 0.75 parasites/ μ l of blood. *Hypostomus regani* and *Hypostomus strigaticeps* showed a significant difference in prevalence. The average densities of parasites were not significantly different among the three fish species. Similar findings were observed for the monthly variations in densities. The parasites found in the three species of catfish studied showed similar morphological characteristics. The morphological data and the statistical analyses used in this study didn't show the formation of groups. The analyses provided evidence of the presence of pleomorphisms in the trypanosomes found in the three studied fish.

Keywords: *Trypanosoma*, fish, *Hypostomus*, catfish, protozoa.

Resumo

Infecções por tripanossomas foram descritas em diversas espécies de peixes, sendo a maioria das descrições baseada nas características morfológicas. Tripanossomas de peixes são heteroxenos e transmitidos por hirudíneos. Este estudo tem como objetivo a avaliação da prevalência e densidade da infecção por *Trypanosoma* sp. no sangue de três espécies de cascudos, *Hypostomus regani*, *H. strigaticeps*, *H. albopunctatus*, do Rio Mogi-Guaçu, Pirassununga, São Paulo, Brasil. Além disso, pretende-se a caracterização das espécies de *Trypanosoma* encontradas no sangue desses peixes, por meio da utilização de técnicas morfológicas e moleculares. A prevalência total de tripanossomas foi de 47,6%, e a densidade média foi de 0,75 parasitas/ μ l de sangue. *Hypostomus regani* e *Hypostomus strigaticeps* apresentaram diferenças significativas em prevalência. As médias das densidades dos parasitas não apresentaram diferenças significativas entre as três espécies de peixes estudadas. O mesmo foi observado com a variação mensal das densidades. Os parasitas encontrados nas três espécies de cascudo possuíam características morfológicas semelhantes. As análises morfológicas e estatísticas obtidas neste estudo não mostraram a formação de grupos. As análises evidenciaram a presença de pleomorfismo dos tripanossomas encontrados nas três espécies de peixes estudados.

Palavras-chave: *Trypanosoma*, peixes, *Hypostomus*, cascudos, protozoários.

*Corresponding author: Marlene Tiduko Ueta. Laboratório de Helmintologia, Departamento de Biologia Animal, Instituto de Biologia, Universidade Estadual de Campinas – UNICAMP, Rua Monteiro Lobato, 255, Bloco J, CEP 13083-862, Campinas, SP, Brasil. e-mail: mtu@unicamp.br

Introduction

Fish trypanosomes have been observed on several continents, and numerous species have been described in salt and freshwater fish (LOPES et al., 1990; EIRAS, 1994).

For fish trypanosomes, the hirudinean aquatic leech is the only known vector host. Fish become infected with trypanosomes during blood feeding by the invertebrate host. There is no evidence of fish infection from the ingestion of infected leeches (BECKER, 1977; LOM, 1979).

Trypanosomes in the blood of infected fish pass through a series of morphological transformations after ingestion by leeches; this transformation process includes the amastigote, espheromastigote, epimastigote, and trypomastigote forms. Soon after the ingestion of infected blood, the flagellates begin to divide in the stomachs of hirudineans (LOM, 1979; LOM & DYKOVA, 1992; EIRAS, 1994; EIRAS et al., 2008).

This sequence of forms is directly related to the level of digestion of the ingested blood, and in most cases, the trypomastigote form appears only when the blood has been fully digested. The hirudinean becomes infectious when the trypanosomes reach the metacyclic trypomastigote form and migrate to the mouthparts of the animal, a process that can last from three to nineteen days (LOM, 1979).

The pathogenicity caused by hirudineans can be more than a simple hemorrhage at the feeding site, where the hirudinean attaches. In severe cases of infections, the leeches can cause anemia and even death of the host (ISLAM & WOO, 1991; WOO, 1995; THATCHER, 2006).

Fish that have recovered from trypanosome infections can develop immunity against re-infection. Thus, the status of the host immune system is an important factor (ISLAM & WOO, 1991; WOO, 1995; OVERATH et al., 1999).

Currently, 59 species belonging to gender *Trypanosoma* have been described in infections of fresh and saltwater fish in Brazil (RIBEIRO et al., 1993). The most significantly affected fish family, with respect to the presence of parasites, is the Loricariidae family, of which 33 species are confirmed to belong to the genus *Hypostomus* and are thus considered to be the most common host (RIBEIRO et al., 1990a). The prevalence of infections in catfish might be due to semi-sedentary habits that enable greater rates of infections by ectoparasites, which in turn transmit the protozoans (FRÓES et al., 1978).

Members of the Loricariidae family are widely distributed and are significantly present in the Paraná basin as well as in the San Francisco basin in Brazil (GARAVELLO & GARAVELLO, 2004; BUCKUP et al., 2007).

The pathogenesis of fish trypanosomiasis is not well known. The parasites live in relative equilibrium with their hosts, which present with low levels of parasitemia. However, changes in the hematological parameters have been observed (BECKER, 1977; EIRAS et al., 1990; EIRAS et al., 2008).

The large variety of hosts, the wide geographic distribution, and the morphometry of parasites represent problems for the descriptions of new species of fish trypanosomes. The identification of new species is also difficult because trypanosomes exhibit polymorphisms (FRÓES et al., 1978), and the taxonomic

identification of fish trypanosome species is furthermore based on differences in the morphological characteristics of the parasites (LOPES et al., 1989; EIRAS et al., 1990).

The traditional taxonomy of *Trypanosoma* species cannot be used as a reliable method for the determination of new species as it is based on morphological differences of trypomastigote forms in the bloodstream. Molecular techniques have been showed as more effective when applied toward the taxonomic determinations to identification of trypanosomes (FIGUEROA et al., 1999; KARLSBAKK & NYLUND, 2006; GU et al., 2007a, b).

Studies of fish trypanosomes can assist in finding solutions to important problems that are common to the studies of the pathogenicity caused by these parasites in mammals. Additionally, such studies will contribute to the phylogeny of the group (LOM, 1979).

The occurrence of parasites in fish under natural conditions can be harmless and might not cause pathogenic alterations; however, in culture conditions the parasitic infections can interfere with the health of the host. Hence, it is important to know which parasites affect fishes species with potential to be farmed, due these infections might cause economical losses of this growing fish farming industry in Brazil.

Objectives

The objectives of this study were to characterize the prevalence and densities of infection by *Trypanosoma* sp. in blood from *Hypostomus albopunctatus*, *Hypostomus regani*, and *Hypostomus strigaticeps*, which were collected from the Mogi Guaçu River, Pirassununga, São Paulo, Brazil; and to characterize morphologically the specimens of *Trypanosoma* that were found in blood from catfish and hirudinean vectors.

Materials and Methods

Monthly collections were performed in the Mogi Guaçu River, Emas Waterfall, in the city of Pirassununga, São Paulo, between February 2008 and February 2009. The fishes, *Hypostomus albopunctatus*, *Hypostomus regani*, and *Hypostomus strigaticeps*, were collected with the aid of nets and were kept in circular 1,000 L tanks at the Center for Research and Management of Continental Fisheries Resources of the Chico Mendes Biodiversity Conservation Institute – CEPTA/ICMBio. The fishes were tagged with numbered plastic tubular labels and biometric measurements were taken before blood collection.

According to the known blood smears technique, blood samples were obtained via cardiac puncture, been obtained five aliquots, each one with 8µl, used to make thick-film smear slides, that were fixed in methanol and stained with Giemsa.

The morphological characteristics of trypomastigote were evaluated and measured from images captured using Leica Image Manager IM 50[®] software. The following parameters were analyzed: body length (BL), free flagella length (FF), distance between the posterior portion and the nucleus (PN), distance between the anterior portion and the nucleus (AN), body width (LC), nucleus width (LN), nucleus length (CN), nucleus area (NRA),

kinetoplast width (LK), kinetoplast length (CK), distance between the kinetoplast and the posterior extremity (DKP), kinetoplast area (ArK), flagellar index (IF), nuclear index (IN), nuclear volume (VN), and kinetoplast volume (VK).

The prevalence and densities were calculated according to the criteria defined by Bush et al. (1997). The Pearson's correlation coefficient were calculated between the densities, weights, and lengths of each fish species. The statistical tests for correlation were performed with a level of significance of 5%.

ANOVA test were performed for each of the morphological variables (BL, FF, PN, AN, LC, LN, CN, Nra, LK, CK, DKP, ArN, IF, IN, VN and VK) using a log transformation in order to achieve normality. For those variables with significant results for the difference of species in the ANOVA tests, the Tukey's multiple comparisons procedure were applied with a combined significance level of 5%.

Multivariate analyses (JOHNSON & WICHERN, 2007) were performed using PCA (principal component analysis) and cluster analysis. Due to high percentage of missing values (> 50%), the variables (FF, IF, DKP) were excluded from the multivariate analysis. In order to get a better approximation to the normal, a log-transformation of all the variables was used. It is also important say that each component is a linear combination of all original variables, in such way that all components are independent.

All statistical analyses were performed using SAS statistical software (SAS Inc., 9.3).

Results and Discussion

Of the 256 collected catfish, 60 were classified as *Hypostomus albopunctatus* (average weight of 288.1g, with standard deviation of 71.4 g; and average length of 28.6 cm, with standard deviation of 2.7 cm), 100 were classified as *Hypostomus regani* (average weight of 322.5 g, with standard deviation of 76.3 g; and average length of 29.6 cm, with standard deviation of 2.5 cm), and 96 were classified as *Hypostomus strigaticeps* (average weight of 281.0 g, with standard deviation of 87.9 g, and average length of 27.3 g, with standard deviation of 3.1 cm). One hundred and twenty two

fishes were positive under laboratory analysis for *Trypanosoma* sp., resulting in an overall prevalence of 47.6% and an overall average density of 0.75 parasites/ μ l of blood (standard deviation 0.75).

In this study, an analysis of *Trypanosoma* sp. infections by individual species of fishes demonstrated the following prevalence and density results: for *Hypostomus albopunctatus*, 46.6% and 0.63 parasites/ μ l of blood (standard deviation = 0.46); for *Hypostomus regani*, 44.0% and 0.78 parasites/ μ l of blood (standard deviation = 0.63); and for *Hypostomus strigaticeps*, 52.0% and 0.78 parasites/ μ l of blood (standard deviation = 0.98).

Two hundred and fifty-five parasites from fishes specimens were found, photographed, and measured in the photomicroscope, as follow: 66 from *H. albopunctatus*, 64 from *H. regani*, and 95 from *H. strigaticeps*. Averages of the measured morphological parameters in the parasites from the three species of fish in the study are showed at Table 1.

The traditional taxonomic identification of fish trypanosome species is based on the morphological characteristics of the parasites. To facilitate morphological description of blood flagellates, morphometrical data were used to divide bloodstream trypanomastigotes into morphotypes. The comparative morphological values of the parasites that were observed in the three studied species and in other fishes species are shown on Table 2. An analysis of variance for each of the variables was performed to verify if there are differences among the three species. The results showed that only for CC and LN there are significant differences (by a 0,05 level Tukey's) between species, where *H. strigaticeps* and *H. albopunctatus* were the same and *H. regani* appear different. Multiple comparisons test were performed and our data set didn't form properly groups. The protozoa measurements obtained from the three species of catfish studied, showed that from the species *H. regani*, *H. strigaticeps* and *H. albopunctatus* were probably belonging to different species (called here *Trypanosoma* morphotype I, *Trypanosoma* morphotype II, *Trypanosoma* morphotype III), this according only considering morphometric analyses showed in Table 1 Lemos et al. (2015) used body length and width to separate the flagellates found in the study in different morphotypes. According to Davies et al. (2005), the nucleus location, which is determined by the distance

Table 1. Average values (in μ m) of the measured parameters in the trypanosomes that were observed in three species of catfish collected between February 2008 and February 2009, in Mogi Guaçu River, Pirassununga, São Paulo, Brazil.

SPECIE	BL	FF	PN	AN	BW	KW	NuW	NuL	NI	Nu-area	DKP	KL	Karea	FI	NV	KV
<i>Hypostomus albopunctatus</i>	29.02	11.50	16.52	10.64	1.96	0.61	1.43	2.85	1.71	4.07	0,45	1.02	0.55	2.90	5.64	0.28
SD	±10.94	±6.24	±5.85	±5.83	±0.72	±0.17	±0.54	±1.06	±0.45	±3.36	±0,14	±0.26	±0.36	±0.89	±6.61	±0.19
<i>Hypostomus regani</i>	24.73	15.45	14.98	8.71	1.94	0.58	1.59	2.96	1.88	4.72	0,33	0.95	0.52	1.95	5.79	0.24
SD	±5.76	±6.06	±3.30	±3.06	±0.45	±0.17	±0.37	±0.65	±0.68	±1.98	±0,19	±0.20	±0.23	±1.15	±3.17	±0.16
<i>Hypostomus strigaticeps</i>	25.01	11.70	15.12	9.43	1.93	0.63	1.50	2.81	1.78	4.08	0,41	0.97	0.65	2.87	5.82	0.29
SD	±7.74	±6.20	±4.57	±4.40	±0.52	±0.21	±0.55	±0.77	±0.61	±3.42	±0,23	±0.24	±0.80	±1.77	±9.03	±0.29

Body length (BL), free flagella length (FF), distance between the posterior portion and the nucleus (PN), distance between the anterior portion and the nucleus (AN), body width (LC), nucleus width (LN), nucleus length (CN), nucleus area (NRA), kinetoplast width (LK), kinetoplast length (CK), distance between the kinetoplast and the posterior extremity (DKP), kinetoplast area (ArK), flagellar index (IF), nuclear index (IN), nuclear volume (VN), and kinetoplast volume (VK). Averages followed by the same letter are not significantly different. SD: standard deviation.

Table 2. Morphometry of the species of *Trypanosoma* sp. as described in different species of fish collected between February 2008 and February 2009, in Mogi Guaçu River, Pirassununga, São Paulo, Brazil. The featured species refer to the present study.

SPECIE	BL	BW	FF	PN	AN	KW	VK	NuW	NuL	IN	VN	HOST	Reference
<i>T. nupelianus</i>	17.60	2.18	12.70	13.77	12.10	0.75	0.28	1.20	3.10	1.14	3.96	<i>Rhinelepis aspera</i>	Eiras et al. (1990)
<i>T. pradoi I</i>	20.20	1.57	8.77	9.63	10.57	0.67	0.67	1.23	2.10	0.91	2.34	<i>Hypostomus ancistroides</i>	Ribeiro et al. (1993)
<i>T. plecostomi I</i>	21.50	2.30	14.00	12.00	-	0.50	-	2.10	2.90	-	-	<i>Hypostomus</i> sp.	Fonseca & Vaz (1928)
<i>T. strigaticeps I</i>	21.90	3.20	12.00	12.50	-	0.60	-	2.10	2.50	-	-	<i>Hypostomus</i> sp.	Fonseca & Vaz (1928)
<i>T. regani II</i>	23.10	1.90	12.30	13.10	-	0.50	-	1.90	3.00	-	-	<i>Hypostomus</i> sp.	Fonseca & Vaz (1928)
<i>T. strigaticeps II</i>	23.50	2.00	12.00	15.00	-	0.50	-	2.00	2.60	-	-	<i>Hypostomus</i> sp.	Fonseca & Vaz (1928)
Trypanosoma Morphotype I	24.73	1.94	15.45	14.98	8.71	0.58	0.24	1.59	2.96	1.88	5.79	<i>H. regani</i>	This study
Trypanosoma Morphotype II	25.01	1.93	11.70	15.12	9.43	0.63	0.29	1.50	2.81	1.78	5.82	<i>H. strigaticeps</i>	This study
<i>T. dominguesi</i>	25.50	2.10	9.40	13.70	12.30	0.70	0.18	1.40	3.90	1.10	7.29	<i>Hypostomus alatus</i>	Lopes et al. (1989)
<i>T. itoi I</i>	25.88	2.18	12.70	13.77	12.10	0.75	0.28	1.20	3.10	1.14	3.96	<i>Hypostomus</i> sp.	Ribeiro et al. (1990b)
Trypanosoma Morphotype III	29.02	1.92	11.50	16.52	10.64	0.61	0.28	1.43	2.85	1.71	5.64	<i>H. albopunctatus</i>	This study
<i>T. lopesi</i>	31.95	2.50	6.40	19.50	4.55	0.70	0.20	1.30	3.45	2.35	7.60	<i>Rhinelepis aspera</i>	Ribeiro et al. (1989)
<i>T. barretoii</i>	36.20	3.87	14.12	19.65	16.65	0.78	0.41	2.65	4.40	1.22	40.37	<i>Hypostomus paulinus</i>	Lopes et al. (1990)
<i>T. itoi II</i>	43.77	3.40	11.17	21.76	22.00	0.84	0.38	1.80	3.57	0.99	8.89	<i>Hypostomus</i> sp.	Ribeiro et al. (1990b)
<i>T. pintoii</i>	45.10	3.30	12.00	25.30	19.80	0.80	0.46	2.20	4.20	1.30	14.40	<i>Hypostomus</i> sp.	Ribeiro et al. (1990b)
<i>T. regani IV</i>	46.10	3.30	16.60	23.30	-	0.90	-	3.10	4.60	-	-	<i>Hypostomus</i> sp.	Fonseca & Vaz (1928)
<i>T. pradoi II</i>	47.58	3.47	17.17	25.62	21.97	0.75	0.75	2.53	4.97	1.17	19.32	<i>Hypostomus ancistroides</i>	Ribeiro et al. (1993)
<i>T. regani V</i>	49.50	2.60	16.00	24.20	-	0.80	-	2.60	4.30	-	-	<i>Hypostomus</i> sp.	Fonseca & Vaz (1928)
<i>T. petenuscii</i>	50.10	3.40	12.80	26.70	23.10	0.80	0.30	3.00	5.00	1.20	30.50	<i>Hypostomus</i> sp.	Carraro et al. (1992)
<i>T. barrosi</i>	53.90	3.30	19.80	30.90	22.70	0.90	0.20	2.80	4.60	1.40	34.90	<i>Hypostomus</i> sp.	Ribeiro et al. (1992)
<i>T. lamanoi</i>	56.70	3.10	13.70	30.30	26.40	1.10	0.30	3.10	5.00	1.10	35.40	<i>Hypostomus ancistroides</i>	Costa et al. (1992)
<i>T. birmanii I</i>	57.20	3.80	9.40	30.30	26.90	0.90	0.43	2.50	4.90	1.10	27.30	<i>Hypostomus commersonii</i>	Ribeiro et al. (1991)
<i>T. affonsoi</i>	57.80	3.60	16.70	30.30	27.50	0.80	0.39	2.80	3.80	1.10	19.60	<i>Hypostomus tietensis</i>	Lopes et al. (1992)
<i>T. birmanii II</i>	59.20	4.50	14.40	31.30	27.80	1.00	0.46	2.60	5.60	1.10	42.40	<i>Hypostomus commersonii</i>	Ribeiro et al. (1991)
<i>T. birmanii III</i>	59.20	4.50	14.40	31.30	27.80	1.00	0.50	2.60	5.60	1.10	42.40	<i>Hypostomus commersonii</i>	Ribeiro et al. (1991)
<i>T. zungaroi IV</i>	61.50	9.00	0.00	32.00	-	7.00	-	4.50	5.50	-	-	<i>Pseudopmelodus zungaro</i>	Fonseca & Vaz (1928)
<i>T. limae</i>	66.64	3.81	11.22	39.58	27.05	1.33	0.42	3.40	5.49	1.46	41.52	<i>Hoplias lacerdae</i>	Lopes et al. (1996)
<i>T. venustissimum II</i>	71.50	2.60	3.50	-	-	1.00	-	-	3.00	1.30	-	<i>Plecostomus plecostomus</i>	Froés et al. (1979)
<i>T. immanis II</i>	114.50	-	6.00	-	-	0.70	-	-	-	1.20	-	<i>Loricariichthys anus</i>	Froés et al. (1978)
<i>T. immanis I</i>	125.60	3.80	7.00	-	-	1.20	-	-	-	1.00	-	<i>Loricariichthys anus</i>	Froés et al. (1978)

Body length (BL), free flagella length (FF), distance between the posterior portion and the nucleus (PN), distance between the anterior portion and the nucleus (AN), body width (LC), nucleus width (LN), nucleus length (CN), nucleus area (NRA), kinetoplast width (LK), kinetoplast length (CK), distance between the kinetoplast and the posterior extremity (DKP), kinetoplast area (ArK), flagellar index (IF), nuclear index (IN), nuclear volume (VN), and kinetoplast volume (VK). Averages followed by the same letter are not significantly different. SD: standard deviation.

between the anterior extremity and the nucleus, is a significant factor for the discrimination of three species of fish trypanosomes reported in Africa (*T. mukasai* (Hoare, 1932); *T. toddi* (Bouet, 1909); *T. tobeyi* (Dias, 1952)). The results shown in the ANOVA described previously were the same as the result obtained in the principal component analysis (PCA). From Figure 1 we can see that with only two components of PCA, 65% of the variability of the data can be explained. The PCAs test showed no grouping of the three species (Figure 2). Multivariate analysis supports no group formation. The results of cluster analyses also showed no formation of groups properly. In the cluster analyses we found 172 groups from our total of 256 elements, which support the findings from the other analyses.

The parasites observed in the three fish species in this study, showed the following morphological characteristics: polymorphisms, dense cytoplasm, clearly visible elliptical nuclei with sharp contours, touching or not the edges of the body and loose chromatin, and terminal and clear kinetoplasts that were located at an average distance of 0.40 μm (\pm 0.23) from the posterior extremities. The bodies gradually tapered toward both ends. The undulating membranes were evident in majority of evaluated specimens, with variation of number of displayed undulations. Some parasites presented with free, long flagella at the anterior extremity, whereas others did not present free flagella. Two morphologically distinct forms were observed; some parasites presented with a slender and short body shape, while others presented with a long and wide body shape and diffuse cytoplasm (Figures 3-5). Occurrence frequency was established according to the variations in body length to assist in the analysis of pleomorphism (Figure 6).

Eigenvalues of the Correlation Matrix				
	Eigenvalue	Difference	Proportion	Cumulative
1	5.32293369	2.18304055	0.4095	0.4095
2	3.13989313	1.67696976	0.2415	0.6510

Figure 1. Eigenvalues of the correlation matrix of the principal component procedure showing the percentage of explanation of the data variability.

The *Trypanosoma* morphotype I, II and III prevalence values observed in this study are greater than those reported by Khan (1977) for North Atlantic fish, and Bureson (2007) for fish from the Pacific. These values are below those reported by Zintl et al. (1997) for European eels and are similar to those reported by Davies et al. (2005) for freshwater fish from Africa.

The density values observed in this study were above those reported by Khan (1977) for *Hippoglossoides platessoides* and

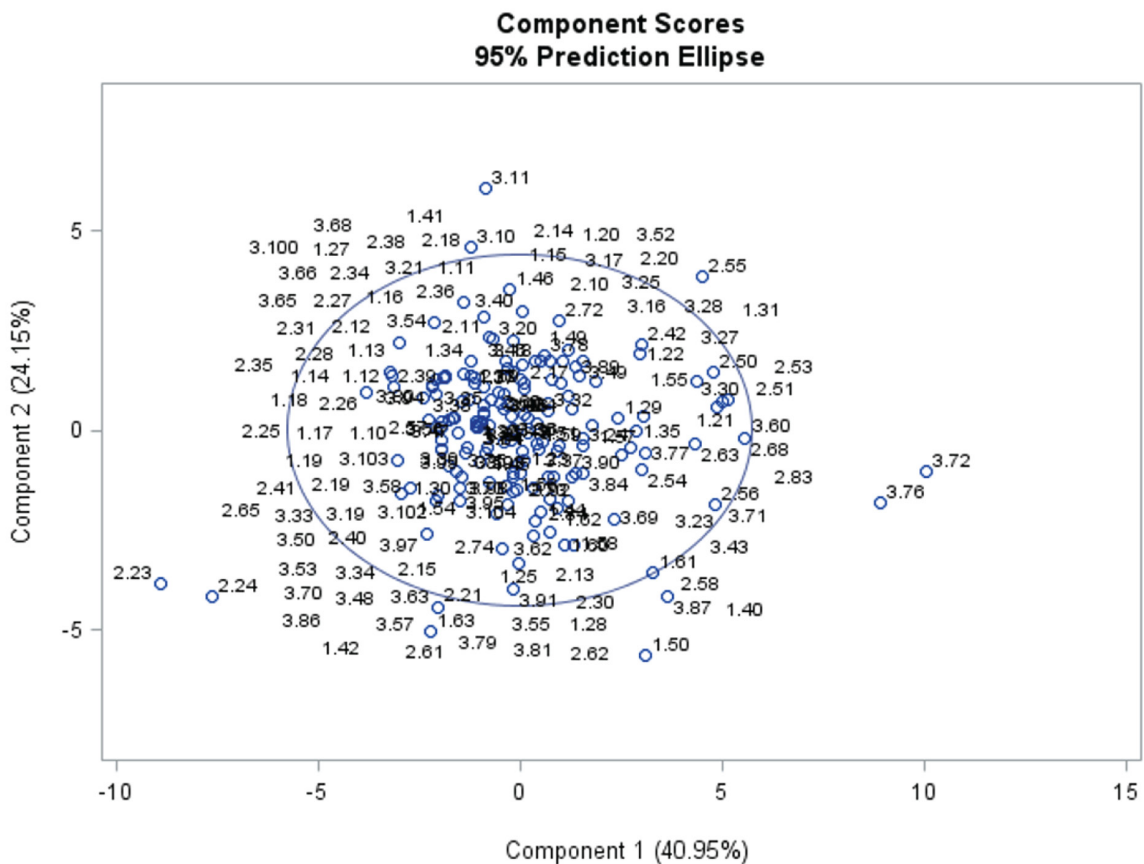


Figure 2. Principal component analyses of the measured parameters in the trypanosomes that were observed in three species of catfish collected between February 2008 and February 2009, in Mogi Guaçu River, Pirassununga, São Paulo, Brazil. (*Hyposyomus regani* called here (1), *Hyposyomus strigaticeps* called here (2) and *Hyposyomus albopunctatus* called here (3)).

Limanda ferruginea, and below those described by Gu et al. (2007b) for *Cyprinus carpio*.

The number of studies that describes *Trypanosoma* species in Loricariidae family is extensive; however, no reference has been reporting the prevalence or densities of these protozoa.

The *Trypanosoma* morphotype I, II and III using just the morphological data can be considered to different species or can be a pleomorphism of the same species. The statistical analyses couldn't give an answer about the possibility of this flagellates showing a grouping of species. By definition, pleomorphism is the presence

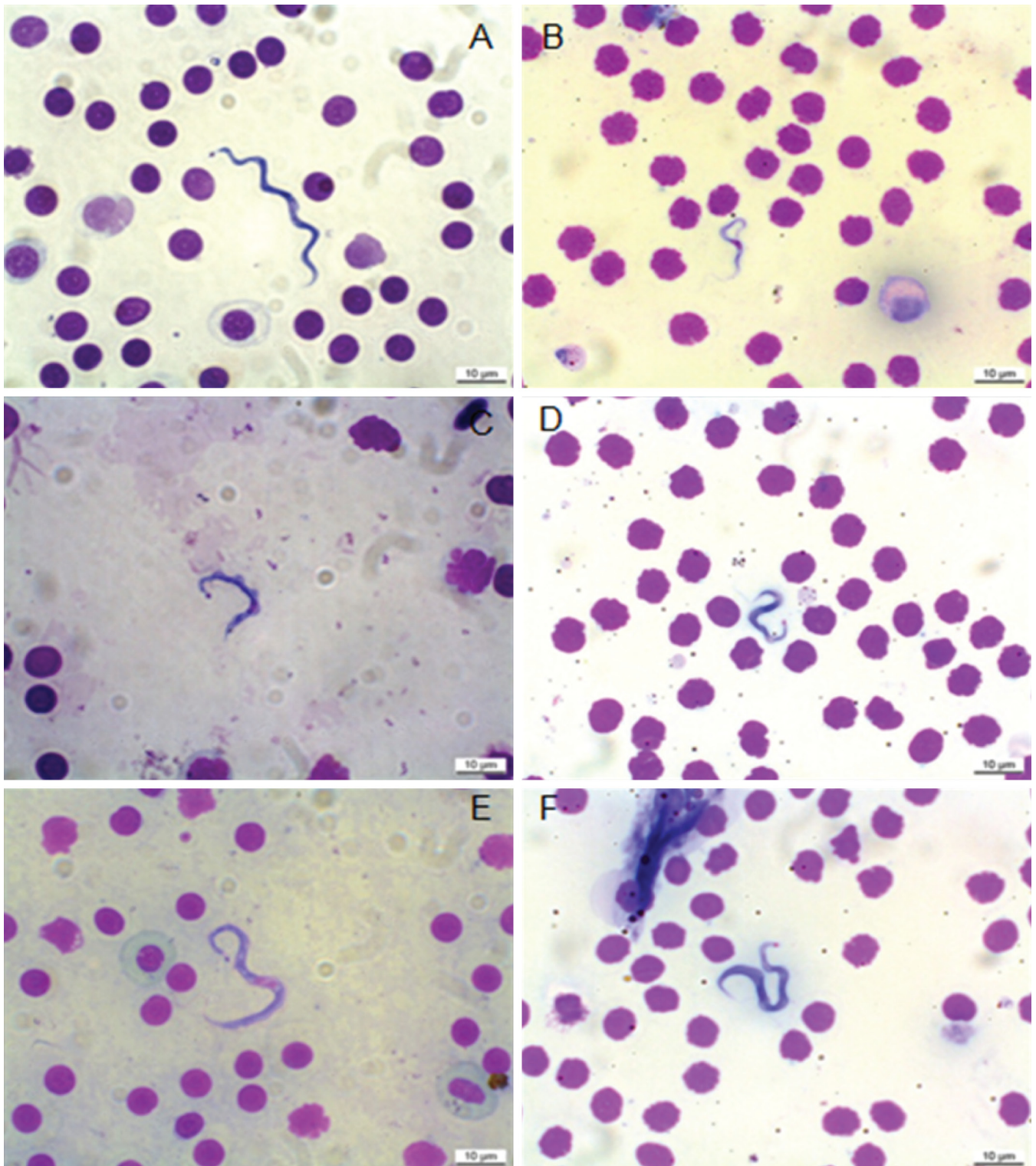


Figure 3. Trypomastigote forms present in the blood of *Hypostomus albopunctatus*. Amplification 100X. A: Long form; B: Short form; C: Membrane with multiple undulations; D: Membrane with few undulations; E: Specimen with free flagella; F: Specimen without free flagella.

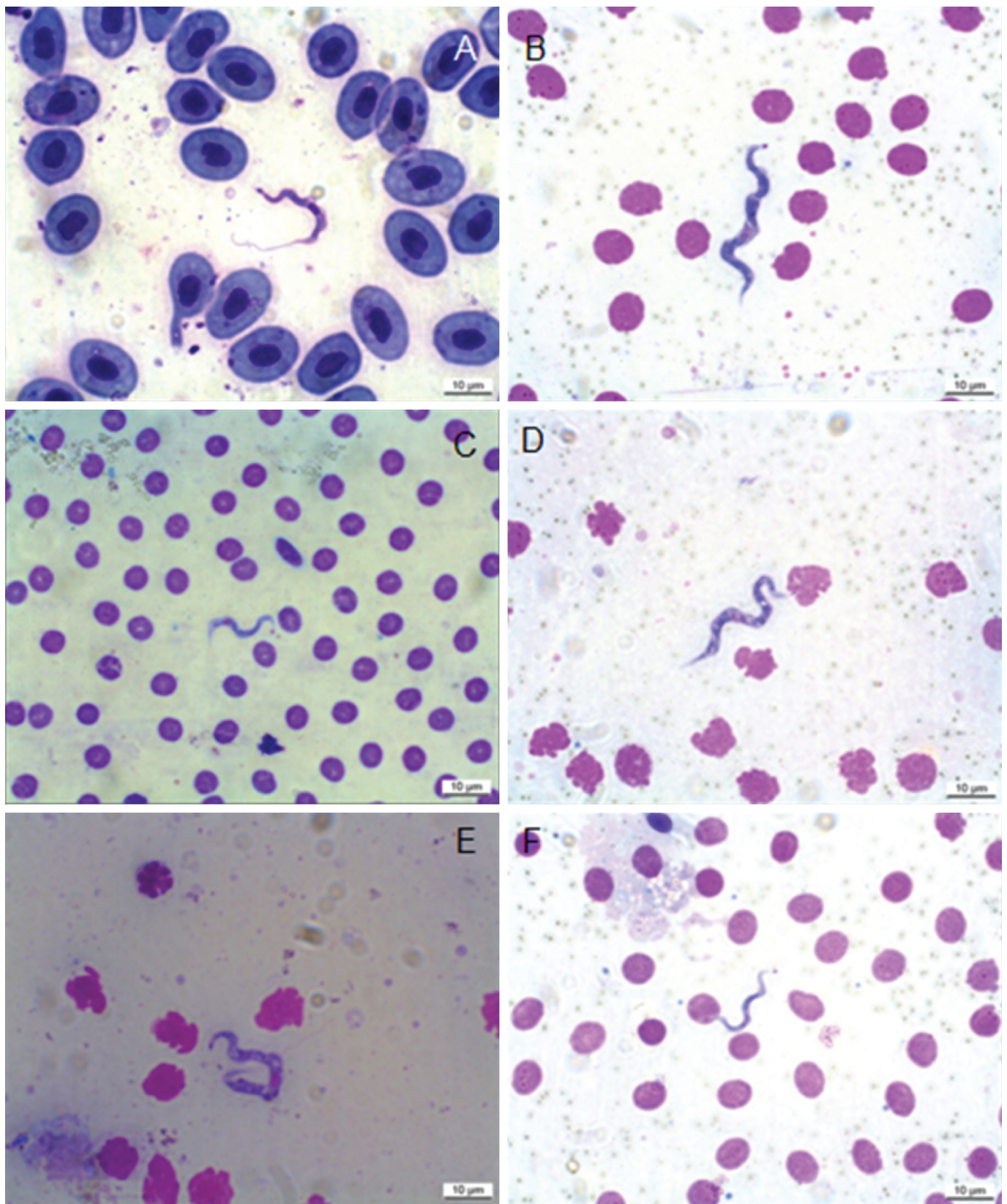


Figure 4. Trypomastigote forms present in the blood of *Hypostomus regani*. Amplification 100X. A: Specimen with free flagella and membrane with multiple undulations; B: Specimen without free flagella and membrane with few undulations; C: Short form; D: Long form; E: Wide form and with diffuse cytoplasm; F: Slender form and with regular cytoplasm.

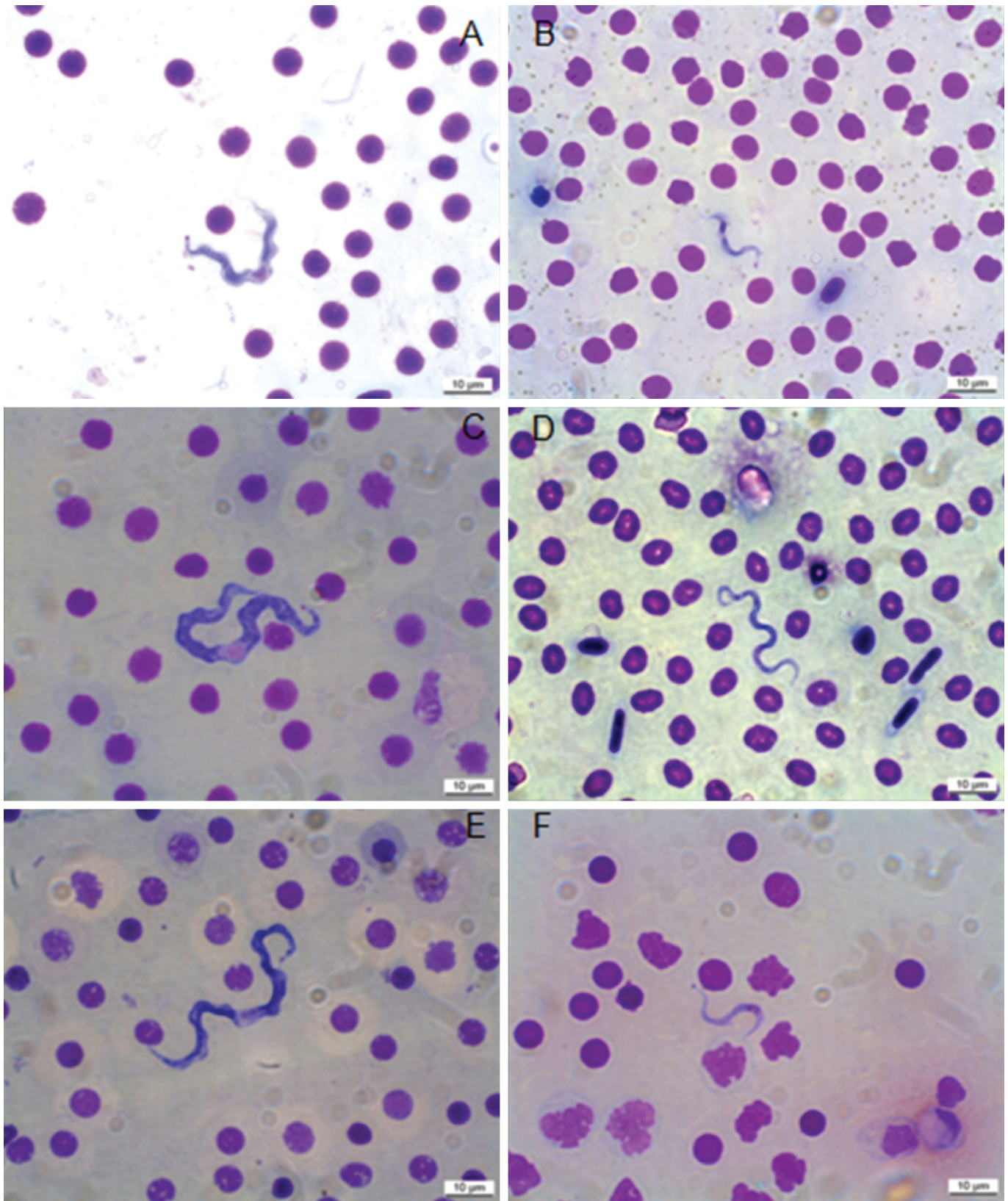


Figure 5. Trypomastigote forms present in the blood of *Hypostomus strigaticeps*. Amplification 100X. A: Specimen with free flagella and membrane with multiple undulations; B: Specimen without free flagella and membrane with few undulations; C: Wide form and with diffuse cytoplasm; D: Slender form and with regular cytoplasm; E: Long form; F: Short form.

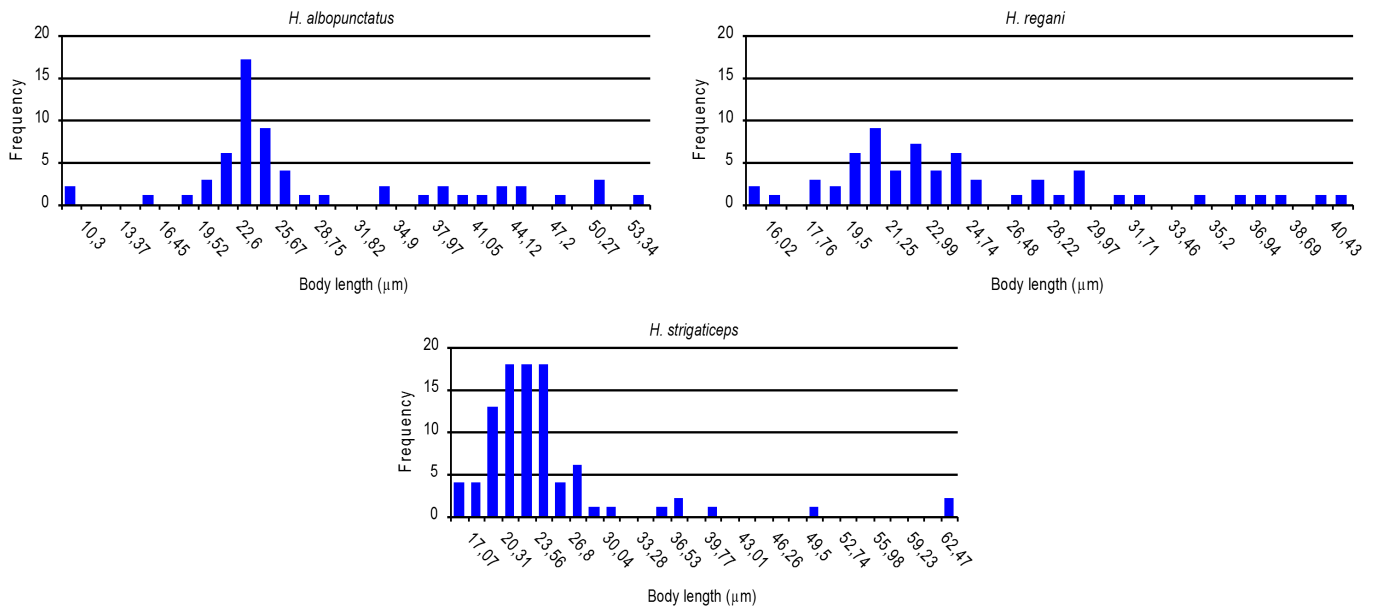


Figure 6. Frequencies distribution of body lengths of trypanosomes that were found in the three studied species of catfish collected between February 2008 and February 2009 at the Mogi Guaçu River, Emas Waterfall, city of Pirassununga, SP.

of various forms of development of a parasite within one host. In blood flagellates from fish, as in other organisms, pleomorphism is understood as a sequential phenotypical manifestation of a genotype that appears in the trypomastigote stage in vertebrate hosts (FERMINO et al., 2015). The pleomorphism can be expressed in the following ways: changes in the total length and width of the body (slender or wider shapes), the number of ripples in the undulating membrane, the presence or absence of a distinct karyosome, the presence of stretch marks on the body surface, the presence and number of cytoplasmic granules, the distance between the kinetoplast and the posterior portion, and the length of the free flagella (LOM, 1979).

According to Gibson et al. (2005), the trypanosomes from freshwater fish found in Europe can be grouped as monomorphic or polymorphic. In the first group, they grouped the parasites showing similar morphologies in the early, acute, and chronic stages of infection, differing only in body length. In the second group, were put the parasites that also showed alterations in length but concomitantly showed variations in body width in the chronic phase of infection.

In a study about the life cycle of *Trypanosoma cobitis*, Letch (1980) reports that the majority of infected fishes that were captured in natural environments were found to have trypanosomes in the peripheral blood, displayed a wide range of morphologies, which may have been the result of multiple infectious events.

The body length variations that were observed in the analyzed trypanosomes specimens, suggest that the pleomorphism appears to be the way to characterize an already established infection. Some authors state that, during the course of infection, the short and slender forms appear at the very beginning, becoming long and wide with the infection maturation (LOM, 1979; LETCH, 1980). Other authors affirm that the pleomorphism is related to the environmental temperatures, with the short and slender forms

appearing during the hottest periods, and long and wide forms appearing during cold periods. In the present study, the differences in the forms of trypanosomes that were observed throughout the year were not characterized as related to environmental temperature or age infection. (WOO, 1994).

Morphologically, the *Trypanosoma* morphotype II and III showed body-length values greater than the few species already describes in the same gender host. In these specimens also was observed that the distance between the front portion and the nucleus was greater than other species (Table 2), including the other sample characterized as *Trypanosoma* morphotype I on this study.

The other *Trypanosoma* morphotype I, described on this study showed body length values (24.89 μm) that were smaller than the most already described by others authors (Table 2). The most curious was that this *Trypanosoma* sp. showed values for all analyzed parameters significantly smaller when compared with *T. regani* IV and *T. regani* V, even when the species were found to infect the same host.

Conclusions

The statistical analyses used in this study didn't show the formation of groups. The flagellates observed in this study provided evidence for the occurrence of pleomorphism in the trypanosomes that were found in the three species of fishes here studied.

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