

Cytogenetic characterization of *Crenicichla* (Pisces, Perciformes, Cichlidae) of the Iguaçú River

S.M.H.K. Mizoguchi, A.L.B. Portela-Castro and I.C. Martins-Santos

Departamento de Biologia Celular e Genética,
Universidade Estadual de Maringá, Maringá, PR, Brasil

Corresponding author: I.C. Martins-Santos
E-mail: icmdsantos@uem.br

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ABSTRACT. Three populations of the genus *Crenicichla*, namely *Crenicichla iguassuensis*, *Crenicichla* sp 1 and *Crenicichla* sp 2, from the Iguaçú River, were analyzed cytogenetically, and their nucleolus organizer regions, constitutive heterochromatin distribution and chromomycin A₃ markings were studied. Karyotype analyses showed a diploid number of 48 chromosomes, made up of 2 metacentric pairs, 2 submetacentric pairs, 7 subtelocentric pairs, and 13 acrocentric pairs for the three *Crenicichla* species and no sexual chromosome differentiation. Nucleolus organizer regions showed strong interstitial marking on the first chromosome pair, coincident with a constriction presented by Giemsa and positive marking by chromomycin. Although

constitutive heterochromatin patterns were also similar, with pericentromeric markings, small differences in the three species could be observed. *Crenicichla* sp 2 presented some chromosomes with bitelomeric markings absent in *Crenicichla iguassuensis* and *Crenicichla* sp 1.

Key words: Karyotype, Nucleolus organizer region banding, C banding, Cichlidae, *Crenicichla iguassuensis*, Perciformes

INTRODUCTION

From an ecological point of view, the Cichlidae family are highly diversified. They belong to the ichthyofauna of Africa, Madagascar, Central and South America, Mexico, southern India, and Sri Lanka (Kullander and Nijssen, 1989; Kullander, 1998). In spite of the group's wide geographic distribution, endemic species may be found in the great African lakes (Kullander, 1998). According to Garavello et al. (1997), pronounced ichthyofauna endemism in the Iguaçú River is due to the Iguaçú waterfalls, which geographically isolate the region from the Paraná River.

According to Nelson (1994), there are 35 genera and approximately 300 species of cichlids in South America, especially in the Amazon basin. More recent studies, however, have given a total of 52 species, distributed into 14 families, including the 36 already mentioned (Garavello et al., 1997). Sixteen of the 23 families of fish reported in the Paraná River are absent in the Iguaçú River, the most important being the anostomids, serrasalmids, curimatids, doradids, and prochilodontids (Agostinho et al., 1997).

Native species, such as *Geophagus brasiliensis*, *Crenicichla iguassuensis* and *Cichlasoma facetum*, and introduced ones, such as *Tilapia rendallii* are among the cichlids of the Iguaçú River (Garavello et al., 1997). Described by Haseman in 1911, *C. iguassuensis* has teeth in several series, depressible posterior teeth, normal lips, upper lateral line distant from the dorsal one, and a dark, elongated body. On the other hand, *Crenicichla* sp, studied by Kullander SO, Pavanelli CS, Lucena CAS and Garavello JC (personal communication), has turgid lips with adipose seams. Lucena and Kullander (1992) analyzed several *Crenicichla* species of the Uruguay River and described *C. tendybaguassu* as having big lips similar to *Crenicichla* sp of the Iguaçú River. A cytogenetic study of the Cichlidae has shown conservative karyotype evolution with regard to diploid number, and all the species of the *Crenicichla* genus studied until the moment present a diploid number of $2n = 48$ chromosomes, as do most species of this family. Enzymatic analysis of 27 loci from *Crenicichla* sp and *C. iguassuensis*, failed to find any diagnostic locus that would indicate two distinct species (Renesto et al., 2001).

Also, the existence of a third species in the same basin has been proposed. Since this species has intermediate-size lips, its presence would suggest three *C. iguassuensis* morphotypes in the Iguaçú River. Our objective was to characterize cytogenetically the different forms and help inform about the hypothesis of more than one *Crenicichla* species in this basin.

MATERIAL AND METHODS

Cytogenetic analyses were undertaken of 71 individuals (24 females and 20 males of *C. iguassuensis*, 4 females and 6 males of *Crenicichla* sp 1, and 8 females and 9 males of *Crenicichla* sp 2), from the Iguaçu River close to the Salto Caxias Reservoir (25° 32' 07 S/53° 28' 59 W) in the State of Paraná, Brazil. Mitotic chromosome preparations were obtained from kidney cells by the air-drying technique described by Bertollo et al. (1978). Constitutive heterochromatin distribution and chromomycin A₃ (CMA₃) were analyzed according to basic procedures suggested by Sumner (1972) and Schweizer (1976), respectively. The nucleolus organizer regions (NORs) were analyzed by silver nitrate (Ag-NOR) staining, following the method described by Howell and Black (1980). Chromosomal types were identified by the arm-ratio criterion proposed by Levan et al. (1964).

RESULTS

Cytogenetic analysis of *Crenicichla iguassuensis* and its morphotypes, *Crenicichla* sp 1 and *Crenicichla* sp 2 showed similar karyotypes with the same diploid number of 48 chromosomes, distributed as 2 metacentric (M) pairs, 2 submetacentric (SM) pairs, 7 subtelocentric pairs, and 13 acrocentric pairs and fundamental number = 56 (Figure 1). Detection of NORs in the three forms by silver nitrate impregnation indicated strong marking in the interstitial region in the short arm of the first SM pair (Figure 1), coinciding with the secondary constriction by Giemsa conventional analysis and with marking by CMA₃ (Figure 2D).

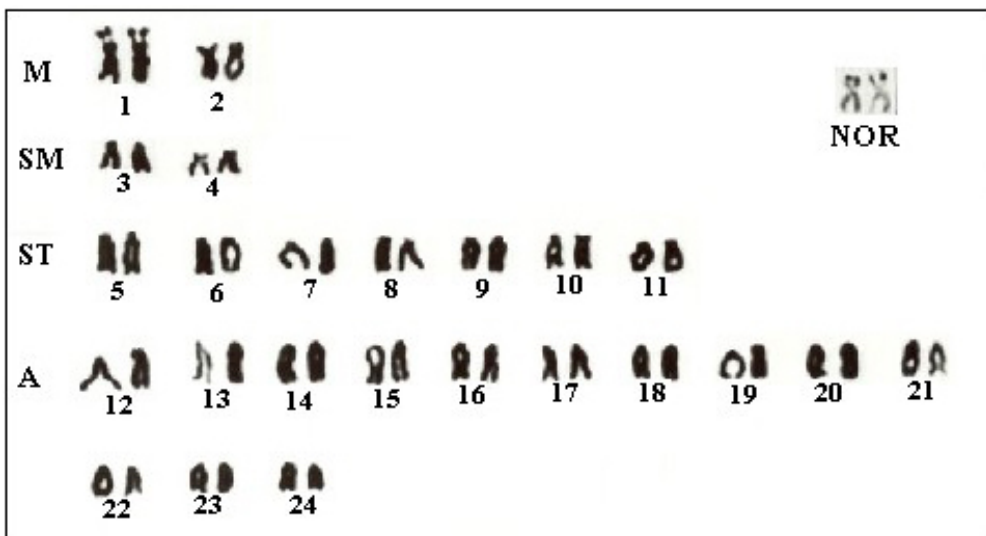


Figure 1. Karyotype of *Crenicichla iguassuensis*, *Crenicichla* sp 1 and *Crenicichla* sp 2. The silver nitrate-nucleolus organizer region (Ag-NOR) chromosome pair is evident. M = metacentric; SM = submetacentric; ST = subtelocentric; A = acrocentric.

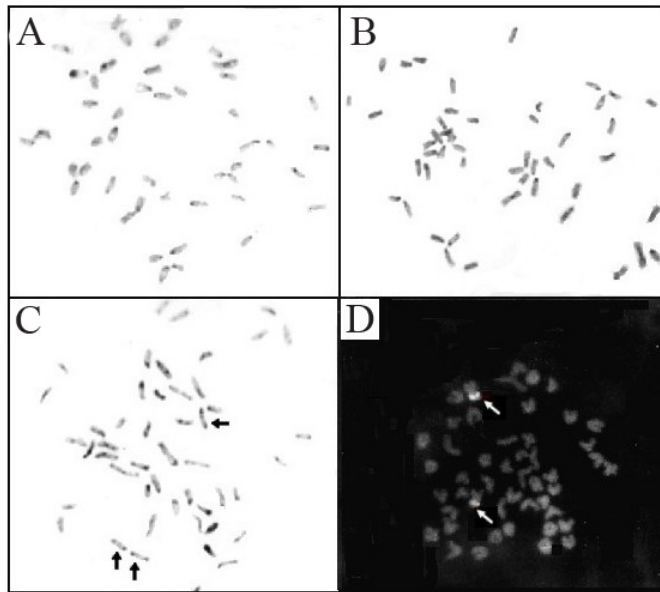


Figure 2. Somatic metaphase of *Crenicichla iguassuensis* (A) *Crenicichla* sp 1 (B) and *Crenicichla* sp 2 (C) evidencing the distribution of the constitutive heterochromatin pattern. Bidelomeric C band in *Crenicichla* sp 2 (arrows). Somatic metaphase of *C. iguassuensis* stained with chromomycin A₃ (CMA₃) representative of the three species (D). Ag-NOR-positive CMA₃ band (arrows).

Constitutive heterochromatin pattern was also similar in the three morphotypes, with pericentromeric markings for most chromosomes and with weak telomeric bands for some others (Figure 2A-C). However, *Crenicichla* sp 2 presented certain chromosomes with bidelomeric markings (Figure 2C). NOR had a negative C band (Figure 2A-C).

DISCUSSION

C. iguassuensis and its two *Crenicichla* sp forms of the Iguaçú River (*Crenicichla* sp 1 and *Crenicichla* sp 2) showed the same diploid number and the same karyotype formula and NOR banding. Although similar, the distribution of constitutive heterochromatin in pericentromeric and telomeric regions was not identical in the three morphotypes. *Crenicichla* sp 2 presented bidelomeric markings, which were absent in the other morphotypes. Analyses suggested that there were no differences that might characterize them as taxonomically different. Renesto et al. (2001) analyzed 27 enzymatic loci in two of the three *C. iguassuensis* forms and found a genetic identity of 0.993, coupled with a lack of a diagnostic locus that would differentiate the two *Crenicichla* morphotypes.

Cichlid cytogenetics has shown a conservative karyotype evolution with regard to diploid number; most of the species present $2n = 48$ chromosomes and many subtelo-acrocentric chromosomes (Thompson, 1979; Feldberg and Bertollo, 1985a; Martins et al., 1995; Loureiro and Dias, 1999). However, their karyotype formulas show variations, in spite of a constant diploid number (Oyhenart-Perera et al., 1975; Feldberg and Bertollo, 1985a; Martins et al., 1995).

Crenicichla species analyzed up to now and other cichlids have also shown a highly conservative chromosome evolution, with a diploid number $2n = 48$ chromosomes, and small

differences in their karyotype formula (Table 1). This fact may be observed through an analysis of the different species in the group with the same karyotype formula (8 M/SM in *Crenicichla* sp B; Salgado et al., 1994, and *Crenicichla* sp, Loureiro and Dias, 1999; 6 M/SM in several species, Table 1). The differences of 6 or 8 M/SM may be a mistake in measurement or differences of chromosome condensation during metaphases under analysis. However, structural rearrangements during the evolutionary process are still a possibility. Consequently, similar karyotypic data in *C. iguassuensis*, *Crenicichla* sp 1 and *Crenicichla* sp 2 corroborate the group's cytogenetic data and show that morphological differences were not visible at the level of the macrostructure karyotype for differentiating the three *Crenicichla* morphotypes of the Iguaçú River.

Table 1. Karyotypic data in the *Crenicichla* genus.

Species	Locality	2n	Karyotype	FN	NOR	Reference
<i>Crenicichla iguassuensis</i>	Iguaçu River	48	8 M/SM + 40 ST/A	56	2	Present study
<i>Crenicichla lacustris</i>	Registro River	48	6 M/SM + 42 ST/A	54	2	Feldberg and Bertollo, 1985a,b
<i>Crenicichla lacustris</i>	Cação River	48	10 M/SM + 38 ST/A	58	2	Brum et al., 2002
<i>Crenicichla lepidota</i>	Miranda River	48	6 M/SM + 42 ST/A	54	2	Feldberg and Bertollo, 1985a,b
<i>Crenicichla lepidota</i>	-	48	6 M/SM + 42 ST/A	54	-	Thompson, 1979
<i>Crenicichla lepidota</i>	Paraná River	48	6 M/SM + 42 ST/A	54	4	Martins et al., 1995
<i>Crenicichla niederleinii</i>	Tibaji River	48	10 M/SM + 38 ST/A	58	2	Loureiro and Dias, 1999
<i>Crenicichla niederleinii</i>	Paraná River	48	12 M/SM + 34 ST/A	62	2	Martins et al., 1995
<i>Crenicichla notophthalmus</i>	-	48	6 M/SM + 42 ST/A	54	-	Thompson, 1979
<i>Crenicichla lucius</i>	-	48	-	-	-	Thompson, 1979
<i>Crenicichla reticulata</i>	Uatumã River	48	6 M/SM + 42 ST/A	54	2	Alves et al., 1999
<i>Crenicichla semifasciata</i>	Miranda River	48	6 M/SM + 42 ST/A	54	2	Feldberg and Bertollo, 1985a,b
<i>Crenicichla</i> sp	Paraná River	48	6 M/SM + 42 ST/A	54	2	Roncati et al., 1996
<i>Crenicichla</i> sp	Itajaí-Açu River	48	8 M/SM + 40 ST/A	56	2	Loureiro and Dias, 1999
<i>Crenicichla</i> sp 1	Iguaçu River	48	8 M/SM + 40 ST/A	56	2	Present study
<i>Crenicichla</i> sp 2	Iguaçu River	48	8 M/SM + 40 ST/A	56	2	Present study
<i>Crenicichla</i> sp A	Amazonas River	48	6 M/SM + 42 ST/A	54	2	Salgado et al., 1994
<i>Crenicichla</i> sp B	Amazonas River	48	8 M/SM + 40 ST/A	56	2	Salgado et al., 1994
<i>Crenicichla strigata</i>	comercial	48	6 M/SM + 42 ST/A	54	-	Thompson, 1979
<i>Crenicichla inpa</i>	Amazonas River	48	6 M/SM + 42 ST/A	54	2	Benzaquem et al., 2002
<i>Crenicichla reticulata</i>	Amazonas River	48	6 M/SM + 42 ST/A	54	2	Benzaquem et al., 2002
<i>Crenicichla cincta</i>	Amazonas River	48	8 M/SM + 40 ST/A	56	2	Benzaquem et al., 2002
<i>Crenicichla vittata</i>	Miranda River	48	6 M/SM + 42 ST/A	54	2	Feldberg and Bertollo, 1985a,b

FN = fundamental number; NOR = nucleolus organizer region; S = metacentric; SM = submetacentric; ST = subtelocentric; A = acrocentric.

A simple NOR has been found in most Cichlids (Feldberg and Bertollo, 1985b; Martins et al., 1995) with variations in sites on the short or long arm of the first chromosome pair of the complement. However, multiple NORs have been reported in *Crenicichla lepidota* (Martins et al., 1995) and in *Cichlasoma paranaense* (Loureiro and Dias, 1998). Sub-terminal secondary constrictions in the first M/SM pair, coinciding with the NOR, have also been reported in most species of the *Crenicichla* genus, indicating this to be the group's ancestral chromosome. A terminal NOR of this chromosome pair has been registered only in *C. cincta* (Benzaquem et al., 2002).

Correspondence between silver nitrate marking and CMA₃ in the NOR site indicating a GC-rich region, has been reported in the three morphotypes analyzed and constitutes a common characteristic among fish species (Wasko et al., 1996; Portela-Castro, 1999). However, it is not a heterochromatic region, a fact that has not been reported for the other species of the group. Although GC-rich NOR is a common characteristic in cichlids and

particularly in the *Crenicichla* genus, NOR-negative C banding seems to be a marker of *Crenicichla* from the Iguaçu River.

Though the distribution of constitutive heterochromatin regions was not identical in the three morphotypes, this was not indicative of different taxonomic unities. Although morphological differences are a consequence of genetic differences, the environment can affect them, hence morphological environmental differences may occur in the absence of reproductive isolation.

Studies on diet and food activity showed that species of the *Crenicichla* genus are piscivorous and that *C. iguassuensis* chiefly ingests fish, decapods (*Aegla* sp) and other invertebrates, whereas *Crenicichla* sp feeds on fish, decapods (*Aegla* sp) and a low percentage of detritus and sediments, not detected in *C. iguassuensis*. Studies on daily food intake show that whereas this species has the highest stomach repletion during the evening-night period, repletion occurs during the day period in *Crenicichla* sp (Hahn et al., 1997).

Diet and feeding activity give important data on species behavior, since feeding source is an important factor in species stabilization and fixing (Agostinho et al., 1997). Although fish with trophic specialization are extant in tropical environments, most have wide feeding flexibility.

In our study, cytogenetic data corroborate those found by isozyme analyses (Renesto et al., 2001), which indicate polymorphisms caused by the feeding habits of the three *C. iguassuensis* forms from the Iguaçu River. However, the group's highly conservative karyotype may be camouflaging real genetic differences that may characterize them as different taxonomic unities. Molecular studies with RAPD, SPAR and mitochondrial DNA techniques have to be undertaken to define, in a satisfactory way, the taxonomic position of the different *Crenicichla* forms found in the Iguaçu River reservoir.

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