



## Cytogenetic divergence between two sympatric species of *Characidium* (Teleostei, Characiformes, Crenuchidae) from the Machado River, Minas Gerais, Brazil

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### Abstract

Cytogenetic studies were performed on two sympatric species of *Characidium*, *C. gomesi* and *C. cf. zebra*, from the Grande River basin, Minas Gerais State, Brazil. Although both species had a chromosome number of 50 with a karyotype exclusively consisting of meta- and submetacentric chromosomes, interspecific diversity was detected concerning the size of the two first chromosome pairs of the karyotypes. Active nucleolus organizer regions (NORs) were located at the terminal position on the long arm of the 17<sup>th</sup> pair of *C. gomesi* and at subterminal position on the long arm of the 23<sup>rd</sup> pair of *C. cf. zebra*. For both species the fluorochrome CMA<sub>3</sub> stained only the NOR-bearing pair of chromosomes. The heterochromatin pattern also showed some differentiation between these species restricted to the centromeric or pericentromeric region of *C. cf. zebra* and practically absent in *C. gomesi*. These data are discussed concerning chromosome diversification in this fish group.

*Key words:* *Characidium gomesi*, *Characidium cf. zebra*, Crenuchidae, NOR-banding, C-banding.

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### Introduction

The genus *Characidium* (family Crenuchidae) is made up of small fish which rarely exceed a standard length of 10 cm and which usually occur in small headwater streams. However, little is known about the chromosomes of the fish in the Crenuchidae despite the fact that this family includes about 80 nominal species (Buckup, 1993a), with the genus *Characidium* alone including 59 nominal taxa which makes it the most diverse genus in the Crenuchidae (Buckup, 1993a, b).

The genus *Characidium* is also the most cytogenetically studied Crenuchidae genera, with cytogenetic data being available on five *Characidium cf. zebra* populations (Miyazawa and Galetti-Jr., 1994; Maistro *et al.*, 1998a; Centofante *et al.*, 2001), one population published as *Characidium cf. fasciatum* and reidentified as *Characidium sp. aff. Characidium zebra*; another as *Characidium cf. lagsantensis* (Miyazawa and Galetti-Jr., 1994); one as *Characidium pterostictum* (Miyazawa and Galetti-Jr., 1994); one as *Characidium sp.* (Miyazawa and Galetti-Jr., 1994); two populations of *Characidium gomesi*,

one published as *C. cf. fasciatum* and reidentified as *Characidium sp. cf. Characidium gomesi* (Maistro *et al.*, 1998a; Centofante *et al.*, 2001; Maistro *et al.*, 2004) and the other as *Characidium lauroi* (Centofante *et al.*, 2003); and one population of *Characidium sp. cf. Characidium alipioi* (Centofante *et al.*, 2003). All these *Characidium* species/populations present a relatively stable karyotypic macrostructure, with a diploid number of  $2n = 50$  and banded chromosomes which are usually meta-submetacentric. Exception to this diploid chromosome number was observed in a Passa-Cinco River (Brazil) *Characidium cf. zebra* specimen which presented one B-chromosome in some cells (Venere *et al.*, 1999), in Quinta and Pardo River (Brazil) *C. gomesi* specimens with 0 to 4 supernumerary chromosomes (Maistro *et al.*, 1998a; 2004) and in a Paiol Grande stream (Brazil) triploid *C. gomesi* specimen (Centofante *et al.*, 2001). Although *Characidium* have demonstrated a constant diploid number of chromosomes, the karyotype structure is variable regarding chromosome shape, sex chromosome systems and number and position of the silver nitrate stained nucleolar organizer region (Ag-NOR) and heterochromatin (Miyazawa and Galetti-Jr., 1994; Maistro *et al.*, 1998a; Centofante *et al.*, 2001; 2003; Maistro *et al.*, 2004).

Since there is no cytogenetic data on *Characidium* from the Minas Gerais region of Brazil, the main objective

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of the present study was to describe the karyotype structure of two sympatric *Characidium* species from the south of Minas Gerais State. The data obtained are discussed concerning some aspects related to the chromosomal evolution of this genus.

## Material and Methods

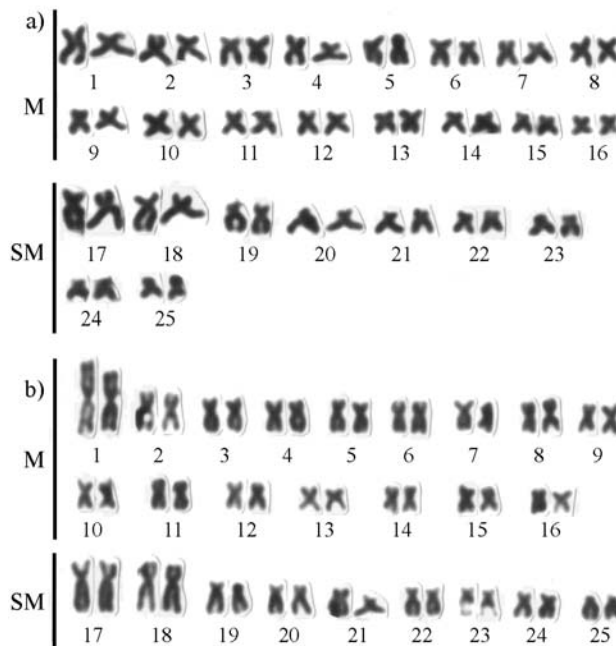
A cytogenetic survey was performed on two sympatric *Characidium* species: 14 female and 6 male ( $n = 20$ ) *Characidium gomesi* and 10 female and 3 male ( $n = 13$ ) *Characidium* cf. *zebra* collected in the Machado River at 22°04.471' S; 46°02.810' W near the town of São João da Mata in the Brazilian State of Minas Gerais. Voucher specimens are deposited at the fish collection of the Brazilian National Museum (Museu Nacional Rio de Janeiro, MNRJ) under catalog numbers MNRJ 28408 for *C. cf. zebra* and MNRJ 28411 for *C. gomesi*.

Mitotic cells were obtained from gill and kidney tissues by the technique described by Foresti *et al.* (1993). Chromosome morphology was determined on the basis of arm ratios as proposed by Levan *et al.* (1964) and the chromosomes were classified as metacentric (M) and submetacentric (SM) and were paired in decreasing order of size. C-banding was performed by the method of Sumner (1972), and silver-staining of the nucleolus organizer regions (Ag-NORs) was performed by the technique of Howell and Black (1980). Chromomycin A<sub>3</sub> (CMA<sub>3</sub>) staining was performed by the method of Schweizer (1980).

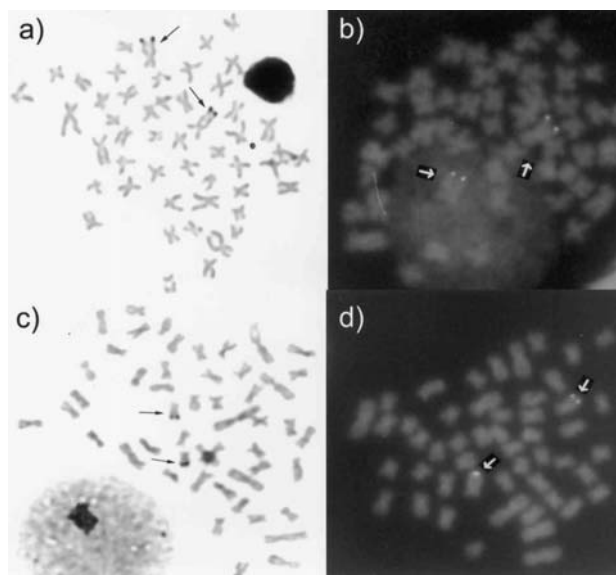
## Results

Giemsa staining showed that the specimens of both *Characidium* species investigated presented the same basic karyotype of  $2n = 50$  (32M + 18SM) with the fundamental number (i.e. the number of chromosomal arms) being equal to 100 (Figures 1a and 1b). However, the first metacentric chromosome pair of *C. cf. zebra* was considerably larger than the second pair while in the *C. gomesi* specimens the first and the second metacentric chromosome pair were similar in size. In *C. cf. zebra* we observed a secondary constriction at the subterminal position on the long arm of the 23<sup>th</sup> chromosome pair (Figure 1b). No chromosome differences were observed between males and females of both species.

The Ag-NOR analysis of both *Characidium* species showed the presence of only one chromosome pair with active NORs. Terminal NORs were observed on the long arm of a pair of major submetacentric chromosomes (pair 17) in *C. gomesi* (Figure 2a) and on the subterminal position on the long arm of the small-sized submetacentric (pair 23), coinciding with the region of secondary constriction in *C. cf. zebra* (Figure 2b). In both species, chromomycin A<sub>3</sub> showed fluorescence only at the NOR sites (Figures 2c and 2d).

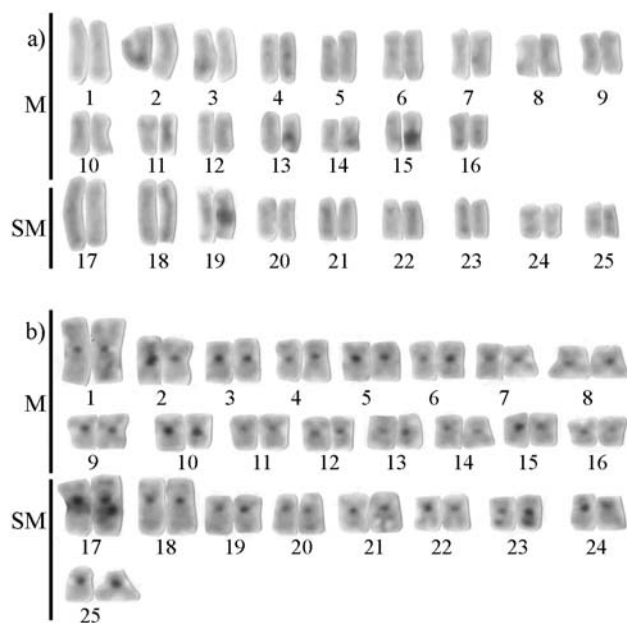


**Figure 1** - Karyotypes of (a) *Characidium gomesi* and (b) *Characidium* cf. *zebra* from the Machado River after conventional Giemsa staining.



**Figure 2** - Somatic metaphases of *Characidium gomesi* (a and b) and *Characidium* cf. *zebra* (c and d) from the Machado River, after silver nitrate (a and c) and CMA<sub>3</sub> staining (b and d). The arrows point NOR-bearing chromosomes.

The C-banding pattern showed heterochromatic blocks at the centromeric or pericentromeric regions of all *Characidium* cf. *zebra* chromosomes and in the subterminal position of the 23<sup>th</sup> chromosome pair (Figure 3a) while *C. gomesi* showed small heterochromatin blocks distributed at the centromeres and telomeres of a few chromosomes, including the telomeric region of the long arm of the 17<sup>th</sup> chromosome pair (Figure 3b).



**Figure 3** - C-banding Karyotypes of (a) *Characidium gomesi* and (b) *Characidium cf. zebra* from the Machado River.

## Discussion

The *Characidium cf. zebra* and *C. gomesi* specimens from the Machado River presented the same diploid chromosome number of  $2n = 50$ , distributed as 32 metacentric and 18 submetacentric chromosomes. This karyotype macrostructure is the same as that observed in the majority of the other *Characidium* species or populations analyzed, with the exception of *C. pterostictum* from the Carlos Botelho Ecological Station in the Brazilian state of São Paulo (Miyazawa and Galetti-Jr., 1994) and *C. lauroi* from the Grande stream, also in São Paulo state, (Centofante *et al.*, 2003) which presented one subtelo-centric chromosome pair and few differences in the number of meta- submetacentric chromosomes.

The *C. cf. zebra* and *C. gomesi* specimens studied by us could be easily separated by the fact that the *C. cf. zebra* first metacentric chromosome pair is larger than the second pair while the *C. gomesi* first metacentric pairs were homogeneous in size. Except for *Characidium. sp. cf. Characidium alipioi* and *C. lauroi* (Centofante *et al.*, 2003) and the *C. gomesi* described in the present paper, the great majority of the *Characidium* species so far studied have a first metacentric pair which is considerably larger than their second metacentric pair. According to Buckup (1993b) *C. zebra* are morphologically primitive and occupy a basal position in the phylogeny of *Characidium*, so the presence of a large first metacentric pair in *Characidium cf. zebra* could be considered primitive for the *Characidium* species that present it, being derived the size homogeneity between the 1<sup>th</sup> and 2<sup>th</sup> metacentric chromosome pairs.

A similar karyotypic pattern has been observed in the cis-Andean genera *Trichomycterus*, made up of small fish that, like *Characidium*, are usually found isolated in the headwaters of small rivers. Sato *et al.* (2004) has shown that the *Trichomycterus* species *florensis*, *reinhardti* and *auroguttatus* form a group in which the first metacentric pair is considerably larger than the second metacentric pair, while *Trichomycterus* sp. aff. *Trichomycterus itatiyae* and *Trichomycterus davisi* form a group for which the first and second metacentric pairs are about the same size, although larger than the other metacentric pairs.

The position of the ribosome sites could clearly differentiate both Machado River populations, *C. cf. zebra* showing Ag-NOR sites in the subterminal position on the long arm of the 23<sup>th</sup> small-sized submetacentric pair while *C. gomesi* presented Ag-NORs at the telomeric position on the long arm of the large submetacentric pair 17. Centofante *et al.* (2003) suggested that species of *Characidium* with a ZZ/ZW sex chromosome systems are more closely related among themselves than to species without such systems. This idea was specially based on the fact that only species of *Characidium* with sex chromosomes presented Ag-NORs in the terminal region of the long arm of a large submetacentric chromosome pair. The Ag-NOR characterization developed in our study showed that the Machado River *C. gomesi* are the only *Characidium* species with Ag-NORs on a large submetacentric chromosome and which do not have a sex chromosome system.

The cytogenetic information available in the literature for other *Characidium* species reports some species with a single chromosome pair bearing Ag-NORs and yet other species with multiple ribosomal sites. Variation in the number of chromosomes bearing Ag-NOR regions is a common feature among characid fishes which has been reported by a number of authors using the Ag-NOR technique (e.g. Almeida-Toledo and Foresti, 1985; Wasko *et al.*, 1996; Jesus and Moreira-Filho, 2003). Maistro *et al.* (1998a) observed multiple Ag-NORs in some Pardo River *Characidium* specimens, however subsequent 18S rDNA FISH analysis of the chromosomes of the 4 specimens showed only one pair bearing ribosomal sites (Maistro *et al.*, 2004). The variable number of ribosomal sites found in the Pardo River population could suggest the occurrence of inter-individual numerical polymorphism of the NOR sites, as has been observed for the *Prochilodus lineatus* chromosome complement (Maistro *et al.*, 2004; Jesus and Moreira-Filho, 2003).

The majority of the *Characidium* species analyzed by us showed Ag-NORs located on small or medium-sized submetacentric chromosomes and, with exception of *C. lauroi*, the Ag-NORs were always located on the long arm. It thus seems that in addition to having a diploid chromosome number of  $2n = 50$  the location of Ag-NORs almost exclusively on the long arm of the chromosomes is another

feature common to *Characidium*. The fact that sympatric *Characidium* species always showed Ag-NORs on different chromosome pairs (Centofante *et al.*, 2001; 2003; present paper) indicates that NOR location is an important cytotaxonomic tool for this group. It may be that FISH analysis with rDNA probes could help in the better understanding of the pattern distribution of ribosome sites on *Characidium* chromosomes.

We found that the Ag-NOR sites were positively stained by the CMA<sub>3</sub> fluorochrome (Figure 2b), suggesting that the rDNA loci of both *Characidium* species studied in may contain spacer sequences or NOR-associated heterochromatin rich in GC base pairs. On the other hand, these data suggest that the C-band positive segments found in both species are not rich in GC base pairs, a characteristic which has been commonly found in several fish species (Amemiya and Gold, 1986; Maistro *et al.*, 2002; Fonseca *et al.*, 2003; among others).

The C-banding pattern could also differentiate the two sympatric *Characidium* species studied by us. In both species heterochromatin preferentially appeared at the centromeres or in the pericentromeric regions but *C. cf. zebra* presented more heterochromatin than *C. gomesi*, which also presented a few telomeric C-band positive blocks. Centromeric and/or pericentromeric heterochromatin has been observed in the majority of the *Characidium* species studied and, in smaller quantities, in *C. gomesi* from the Paiol Grande stream (Centofante *et al.*, 2001) as well as in the *C. sp. cf. C. gomesi* from the Pardo River (published as *C. cf. fasciatum* by Maistro *et al.*, 1998a) and *C. gomesi* studied by us. The available C-banding data shows that the *Characidium* taxa most related to *gomesi* present fewer heterochromatin-bearing chromosomes than other *Characidium* species.

Maistro *et al.* (1998a) described a ZZ/ZW sex-chromosome system for Pardo River *C. gomesi*, where the Z and W chromosome have the same shape and size and are differentiated from each other by the total amount of heterochromatinization of the W chromosome. Centofante *et al.* (2001) found a similar sex chromosome system in *C. gomesi* from the Paiol Grande stream, these fish also presenting high heterochromatinization of the W chromosome but in this case this chromosome was small in comparison to the Z chromosome. Since we detected no sex chromosome differentiation in *C. gomesi* and considering the fact that these fish live in headwaters, show low geographical mobility and form local populations, attributes can facilitate speciation, we feel that *C. gomesi* from the Machado River could represent a new *Characidium* species.

On the basis of the cytogenetic data available on *Characidium* species we suggest that the general trend of this group towards karyotypic diversification is similar to other fish groups that maintain a conserved karyotype macrostructure but that are quite divergent in terms of NOR

location, heterochromatin distribution and the occurrence of sex chromosomes (Koehler *et al.*, 1997; Pereira *et al.*, 2002; for example). The specific characteristics observed in the chromosome structure of the genus *Characidium* are probably due to the fact that this nominal species is composed of isolated populations found in the headwaters of small tributaries, which probably followed different routes of chromosomal diversification. These characteristics make *Characidium* an excellent group for supplementing the evolutionary studies which have been carried out with *Astyanax scabripinnis* and *Trichomycterus* that have similar ecological characteristics (Moreira-Filho and Bertollo, 1991; Maistro *et al.*, 1998b; Borin and Martins-Santos, 1999). Since only a few *Characidium* species have so far been investigated, further studies on other *Characidium* species and populations are necessary in order to better understand the processes underlying chromosome diversification in this group of fish.

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