Comparative cytogenetic studies in species of the subfamily Callichthyinae (Teleostei: Siluriformes: Callichthyidae)

SHIMABUKURO-DIAS CRISTIANE KIOKO¹, CLAUDIO OLIVEIRA^{1,*} and FAUSTO FORESTI¹

Abstract — In the present study, the karyotype of three species (nine populations) of the Callichthyinae subfamily were investigated with the objective of better understanding the pattern of relationship among the genera that compose the subfamily. Among the four populations of *Callichthys callichthys* studied, two showed 2n=56 chromosomes and two 2n=58 chromosomes. Up to eight additional microchromosomes were observed in the sample from Marília. The three populations of *Hoplosternum littorale* displayed the same number of chromosomes, 2n=60, and karyotypic constitution, 6M+2SM+52A. The two populations of *Megalechis personata* showed 2n=62 chromosomes and similar karyotypic formulae, 8M+54A and 6M+2SM+54A. Terminal Ag-NORs were found in one chromosome pair of *C. callichthys*, *H. littorale*, and *M. personata* from Itiquira, and in two pairs in *M. personata* from Rio Branco. The populations of *C. callichthys* showed C-band positive segments in centromeric and pericentromeric position and the populations of *H. littorale* and *M. personata* exhibited C-band positive segments in centromeric and/or interstitial position. Contrarily to the extensive chromosome rearrangements verified in the Corydoradinae subfamily, in the Callichthyinae subfamily a small number of changes seems to have occurred in its karyotypic evolution.

Key words: Ag-NOR, C-band, chromosomes, fish cytogenetic, karyotypic evolution.

INTRODUCTION

The Callichthyidae family is composed by 177 species separated in two subfamilies, Corydoradinae and Callichthvinae (REIS 2003). The Corvdoradinae subfamily is composed by the genus Aspidoras (18 species), Corydoras (143 species), and Brochis (3 species) and the Callichthyinae subfamily is formed by the genera Callichthys (2 species), Dianema (2 species), Hoplosternum (3 species), Lepthoplosternum (4 species), and Megalechis (2 species) (REIS 2003). According to the phylogeny proposed by Reis (1998), based on morphological characters, Dianema and Hoplosternum compose one clade which is the sister group of Megalechis. Lepthoplosternum is the sister group of Megalechis, Dianema, and Hoplosternum and Callichthys is the most primitive genus in the subfamily.

Cytogenetic analysis conduced in the Callichthyinae subfamily showed that the diploid number in this group has a small variation, from

2n=52 to 2n=66 (Porto et al. 1992; Oliveira et al. 1993), when compared to the extensive karyotypic variability found among the Corydoradinae, mainly in the genus Corydoras, which diploid numbers range from 2n=40 in Corydoras nattereri (Oliveira et al. 1990) to 2n=134 in C. aeneus (Turner et al. 1992). This difference in the karyotypic variability between both Callichthyidae subfamilies is positively related to the number of species that constitute each one; thus, while the genus Corydoras is the most speciose among the Siluriformes, the other genera of the Callichthyidae family have a small number of species.

In the present study the karyotype of three species of Callichthyinae are described, after the chromosome staining with Giemsa, the localization of nucleolus organizer regions with Silver Nitrate (Ag-NORs), and C-banding, with the objective of increasing the cytogenetic data in the group, searching for new elements that could be used in order to better understand the pattern of relationships among its genera.

MATERIAL AND METHODS

The cytogenetic studies were done with three species (nine populations) of the Callichthyinae

¹ Departamento de Morfologia, Instituto de Biociências, Universidade Estadual Paulista, 18618-000, Botucatu, SP, Brazil.

^{*} Corresponding author: fax +55 14 3815-3744; e-mail: claudio@ibb.unesp.br.

subfamily. Their taxonomic position, collection site, and number and sex of the analyzed specimens are shown in Table 1. After processing, fishes were identified and stored as vouchers at the fish collection of Laboratório de Biologia de Peixes (LBP), Instituto de Biociências, Universidade Estadual Paulista, Botucatu, São Paulo, Brazil.

Chromosome preparations were carried out according to Foresti *et al.* (1993). Silver-staining of the nucleolar organizer regions was performed according to the technique proposed by Howell and Black (1981), the C-banding was performed as described by Sumner (1972). Chromosome morphology was determined on the basis of arm ratio as proposed by Levan *et al.* (1964) and chromosomes were classified as metacentrics (M), submetacentrics (SM), subtelocentrics (ST) and acrocentrics (A).

RESULTS

Among the four local populations of *C. calli-chthys* studied, two from Marília and Corumbá had 2n=56 chromosomes (Fig. 1) and two from Pindamonhangaba and Embu-Guaçu had 2n=58 chromosomes (Fig. 2). In the population of Mar-

ília, the occurrence of 0 to 8 microchromosomes was observed (Fig. 1a). One specimen from the population of Corumbá exhibited a large additional M chromosome, with size similar to those of the first M pair (Fig. 1c). The three populations of *H. littorale* had the same number of chromosomes (2n=60) and karyotypic constitution (6M+2SM+52A) (Fig. 3a). The two samples of *M. personata* had 2n=62 chromosomes and similar karyotypic formulae, 8M+54A and 6M+2SM+54A, respectively (Figs. 3a and 3b). In the species in which males and females were analyzed, differences between the sexes were not observed.

Single terminal Ag-NORs, located in the short arm of the chromosomes were found in *C. callichthys, H. littorale*, and *M. personata* from Itiquira (Figs. 1 to 3). The species *M. personata* from Rio Branco had terminal Ag-NORs in two M pairs, located in the short arms of the first chromosome pair and in the long arm of the second chromosome pair (Fig. 3c). All populations of *C. callichthys* displayed size polymorphism in the Ag-NORs (Figs. 1 and 2).

The populations of *C. callichthys* showed C-band positive segments in centromeric position in almost all chromosomes (Fig. 4). The microchromosomes of *C. callichthys* from Marília were entirely C-band positives (Fig. 4a). In the speci-

Table 1 — Studied species, collection numbers (LBP), number and sex of the studied fish, collection sites, diploid number (2n) and karyotypic data obtained for the studied species.

Genera and species	Collec- tion number	Number of specimens	Locality	2n	Karyotype	Number of Ag- NORs- bearing pairs	Number and size of super- numerary chromosomes
Callichthys							
C. callichthys	485	1/1	Pombo stream, Marília, São Paulo	56	22M + 16SM + 18ST	1	0-8 micro- chromosomes
C. callichthys	622	4/1	Convento stream, Pindamonhangaba, São Paulo	58	18M + 14SM + 26ST	1	
C. callichthys	848	2/0	Santa Rita stream, Embu-Guaçu, São Paulo	58	18M + 16SM + 24ST	1	
C. callichthys	1555	3/0	Corumbá, Mato Grosso	56/57	20M + 16SM + 20ST	1	1 macro- chromosome
Hoplosternum							
H. littorale	569	2/1	Guaíba river, Eldorado do Sul, Rio Grande do Sul	60	6M+2SM+52A	1	
H. littorale	466	1/3	Jurumirim reservoir, Angatuba, São Paulo	60	6M+2SM+52A	1	
H. littorale	688	1/1	Tributary of the Piraí river, Poconé, Mato Grosso	60	6M+2SM+52A	1	
Megalechis			,				
M. personata	1554	2/0	Tributary of the Itiquira river, Itiquira, Mato Grosso	62	6M+2SM+54A	1	
M. personata	921	1/1	Almoço stream, Rio Branco, Acre	62	8M+54A	2	

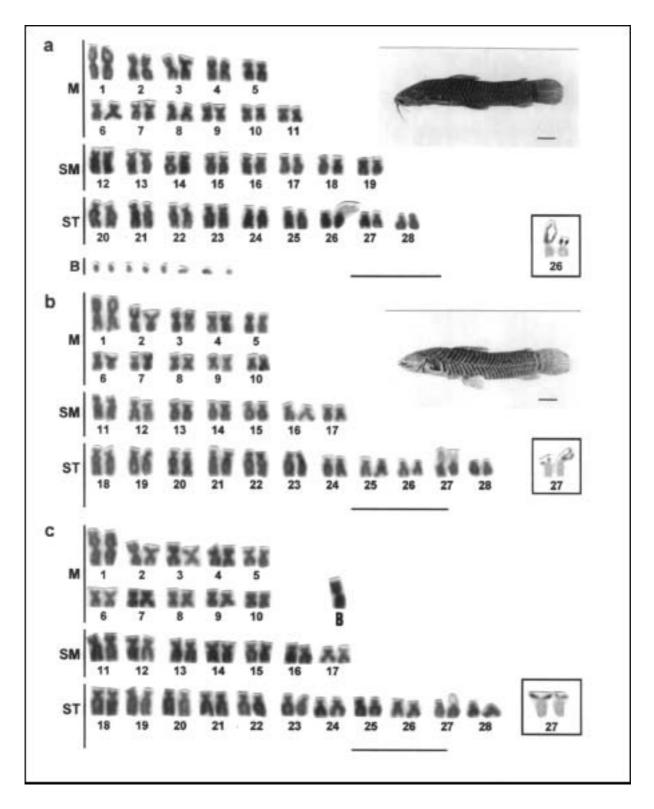


Fig. 1 — Karyotypes, Ag-NORs and specimens of: a) Callichthys callichthys from Marília, 2n=56, with eight additional supernumerary chromosomes (B); b) C. callichthys from Corumbá, 2n=56; and c) C. callichthys from Corumbá, 2n=57, with a putative additional supernumerary chromosome (B). Scale bars: $10 \, \mu m$ (karyotype) and $10 \, mm$ (specimen).

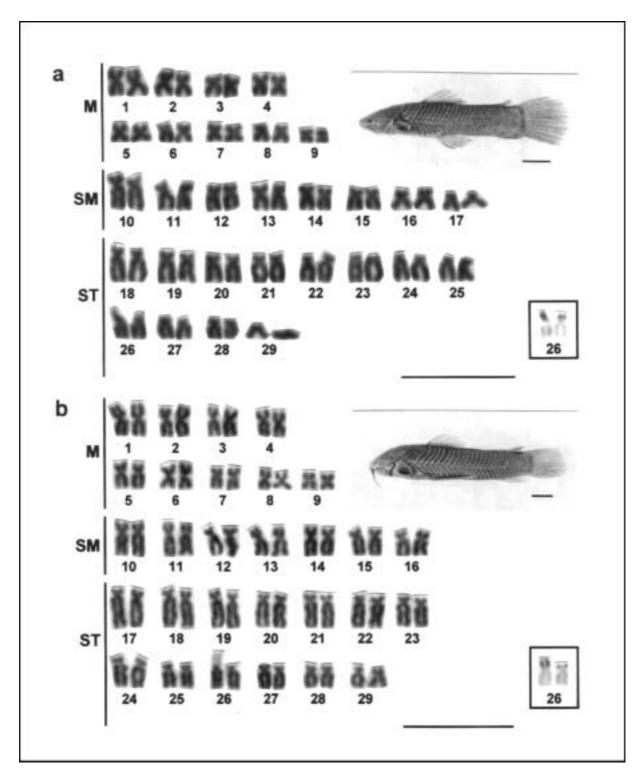


Fig. 2 — Karyotypes, Ag-NORs and specimens of: a) *Callichthys callichthys* from Embu-Guaçu, 2n=58; and b) *C. callichthys* from Pindamonhangaba, 2n=58. Scale bars: 10 μm (karyotype) and 10 mm (specimen).

men of *C. callichthys* from Corumbá with 2n=57 chromosomes, the additional M chromosome exhibited a small C-band positive segment in the centromeric position, and one of the homologues

of the second M pair and of the first ST pair presented the short arms entirely C-band positive (Fig. 4c). The populations of *H. littorale* and the samples of *M. personata* had the C-band positive

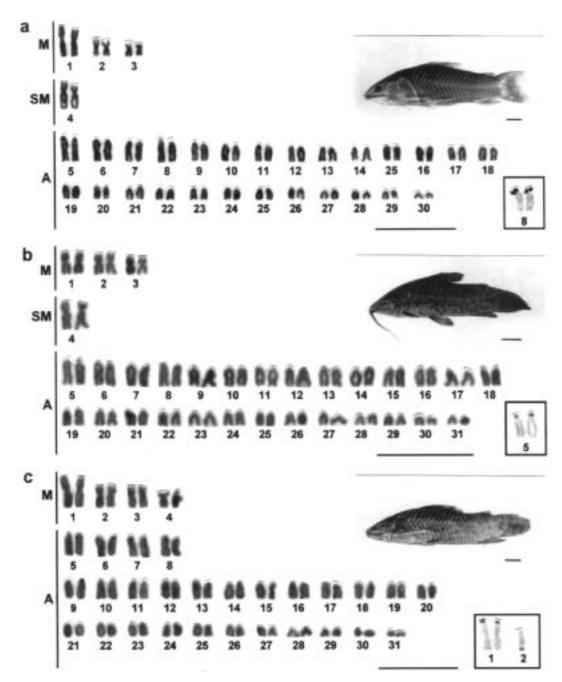


Fig. 3 — Karyotypes, Ag-NORs and specimens of: a) *Hoplosternum littorale* from Poconé, 2n=60; b) *Megalechis personata* from Itiquira, 2n=62; and c) *Megalechis personata* from Rio Branco, 2n=62. Scale bars: $10 \, \mu m$ (karyotype) and $10 \, mm$ (specimen).

segments distributed in centromeric and/or interstitial position in all chromosomes (Fig. 5). The Ag-NORs were C-band positives in all analyzed species (Figs. 4 and 5).

DISCUSSION

The available data for *C. callichthys* showed that this species has a noticeable variation regard-

ing to the diploid numbers, ranging from 2n=52 to 2n=58 chromosomes (Porto *et al.* 1992; Oliveira *et al.* 1993; present study), with the occurrence of several cytotypes in different Brazilian hydrographic basins. Thus, in the Parana-Paraguay basin three cytotypes, with different diploid numbers were found: one with 2n=54, in the Paraná river in the municipality of Porto Rico (I. C. Martins Santos, personal communication);

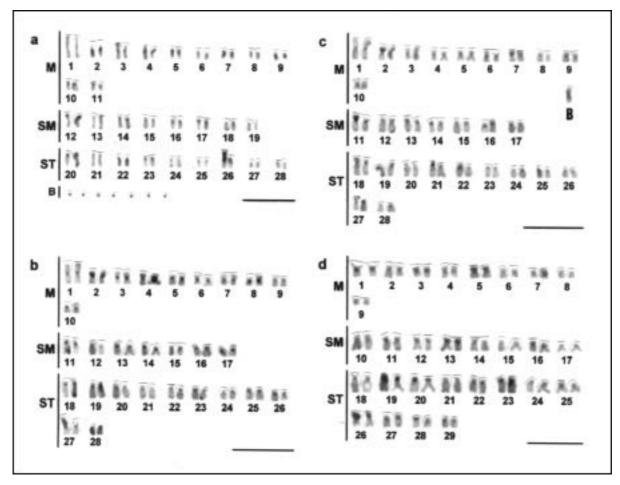


Fig. 4 — C-banded karyotypes. a) *Callichthys callichthys* from Marília, 2n=56, with seven additional supernumerary chromosomes (B); b) *C. callichthys* from Corumbá, 2n=56; c) *C. callichthys* from Corumbá, 2n=57, with a putative additional supernumerary chromosome (B); d) *C. callichthys* from Embu-Guaçu, 2n=58. Scale bars: 10 μm.

one with 2n=56, in the north part of Argentina (Sanchez and Fenocchio 1996), in the Paraguay river in the municipality of Corumbá (present study), and in a tributary of Tietê river in the municipality of Marília (present study); and one with 2n=58 in a tributary of right side of upper Tietê river in the municipality of Guarulhos (Oliveira et al. 1993) and in a tributary of left side of upper Tietê river in the municipality Embu-Guaçu (present study).

While the cytotypes with 2n=56 chromosomes have been found only in the Parana-Paraguay basin, cytotypes with 2n=58 chromosomes have been found in freshwater coastal rivers in the South and Southeast Brazilian regions in the municipality of Tapes and Viamão (CALCAGNOTO 1989), Itanhaém (OLIVEIRA *et al.* 1993), and Pindamonhangaba (present study). Moreover, fish with 2n=58 were also found in the Amazon basin (PORTO *et al.* 1992). In the Amazon basin cyto-

types with 2n=52 and with 2n=54 chromosomes were also found (Porto *et al.* 1992).

From the taxonomic point of view, the genus *Callichthys* have only two valid species: *C. callichthys* and *C. fabricioi*; however, 13 species were already described and now these names are considered junior synonymous of *C. callichthys* (Reis 2003). The presence of different cytotypes in the species *C. callichthys* suggest the occurrence of several unidentified species, reinforcing the proposal of Reis (1998), agreeing that the group needs a general taxonomic review.

Among the neotropical freshwater fish, supernumerary chromosomes, with different sizes, have been found in several families, such as Anostomidae, Characidae, Crenuchidae, Curimatidae, Parodontidae, Prochilodontidae, Callichthyidae, Loricariidae, Pimelodidae, Cichlidae, and Coregonidae (Salvador and Moreira-Filho 1992; Mizoguchi and Martins-Santos 1997; Vênere et al. 1999). In general the supernumerary chro-

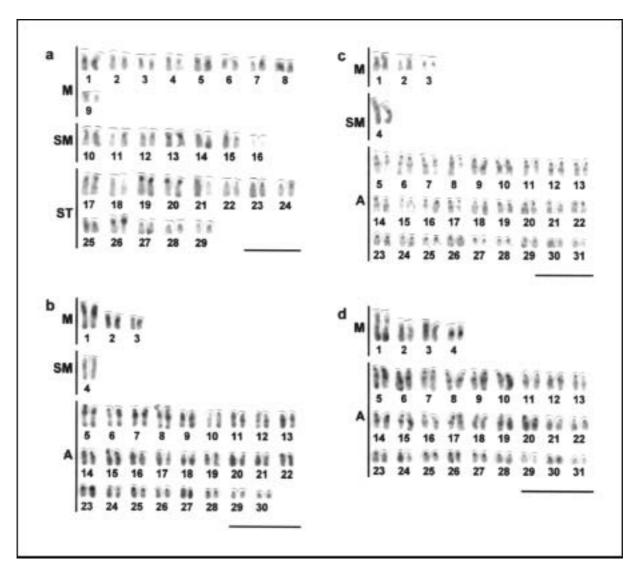


Fig. 5 — C-banded karyotypes. a) *C. callichthys* from Pindamonhangaba, 2n=58; b) *Hoplosternum littorale* from Poconé, 2n=60; c) *Megalechis personata* from Itiquira, 2n=62; and d) *Megalechis personata* from Rio Branco, 2n=62. Scale bars: 10 µm.

mosomes are very small (microchromosomes) and their number range from one to two, with the exception of Moenkhausia sanctafilomenae (Foresti et al. 1989) and C. callichthys (CALCAGNOTO 1989), which are species where up to eight and 16 microchromosomes, respectively were found. In a sample of C. callichthys from Itanhaém the presence of up to three supernumerary microchromosomes were found (OLIVEIRA et al. 1993). In both samples of *C. callichthys* with supernumerary chromosomes previously studied, the specimens had 2n=58. In the present study, an interesting characteristic observed in one specimen from Marília, with 2n=56 chromosomes, was the presence of up to 8 microchromosomes, with the common characteristics found in the supernumerary chromosomes. The existence of a cytotype with 2n=56 and with supernumerary chromosomes is interesting because it shows that, besides de difference in the diploid number in the State of São Paulo, the supernumerary chromosomes in this genus are not restricted to the cytotypes with 2n=58.

The occurrence of supernumerary chromosomes in *C. callichthys* was also described for a population from Argentina, municipality of Reconquista (Sanchez and Fenocchio 1996), in which was observed a diploid number of 2n=56 and from one to two additional small size M. In the Callichthyidae family, small supernumerary chromosomes were also described to *Corydoras aeneus* (Oliveira *et al.* 1988). The analysis of a sample of three specimens of *C. callichthys* from

Corumbá (present study) showed that two had 2n=56 chromosomes and a third one 2n=57, being the additional chromosome a large M. Although this additional chromosome could be a supernumerary chromosome, the analysis of the chromosomes of the fish with 2n=57, after the C band staining, showed the presence of two chromosomes with only one homologue with the short arm C-band positive (pairs 2 and 11 in the Fig. 4c). The presence of this additional chromosome and at least two chromosome pairs without evident homologues permit to suggest that this fish with 2n=57 could be a hybrid between fishes with 2n=56 and 2n=58 chromosomes, since these two cytotypes occur in the Paraná-Paraguay basin. The analysis of more specimens is required to resolve what this additional chromosome and the polymorphism of C-band actually mean.

All populations of *C. callichthys* analyzed in the present study had a single chromosome pair with Ag-NORs and C-band positive segments mainly found in centromeric and pericentromeric positions, as described before (OLIVEIRA *et al.* 1993; SANCHEZ and FENOCCHIO 1996).

The three populations of *Hoplosternum littorale* studied had the same diploid number, karyotypic structure, number and position of Ag-NORs and pattern of distribution of C-band positive segments. These characteristics were also found in two other populations of *H. littorale* and for some specimens identified as *Hoplosternum* sp. (Porto and Feldberg 1992; Oliveira *et al.* 1993). Although, *H. littorale* is the species with wider geographic distribution among the Callichthyidae (Reis 1997), the karyotypic similarity among the populations studied suggest that this species may be introduced in some rivers where it was not found some years ago, such as in the rivers of the upper Parana basin.

The sample of Megalechis personata from Rio Branco had 2n=62 chromosomes and karyotypic structure very similar to those described for a population from the Araguaia river (Vênere 1998). The results obtained for the sample of M. personata from Itiquira were very similar to those obtained for the sample from Rio Branco although this sample was collected in the Parana-Paraguay basin. Additionally, two samples described as *Hoplosternum thoracatum* and a sample described as H. aff. thoracatum, both names considered junior synonymous of Megalechis according to REIS (1997), had 2n=62, 2n=64, and 2n=66, respectively (Porto and Feldberg 1992; Porto et al. 1992). While the populations of M. personata (Vênere 1998; present study) showed many A chromosomes, both samples studied by PORTO and FELDBERG (1992) and PORTO *et al.* (1992) presented several M/SM chromosomes, suggesting the occurrence of several chromosomal rearrangements, as inversions, during the evolutionary history of the group. The presence of different cytotypes may also indicate the occurrence of undescribed species in the genus.

The presence of 2n=56 chromosomes, single terminal NORs, and a small amount of centromeric heterochromatin are characteristics found in Diplomystes mesembrinus, a species that belongs to the most primitive family of the order Siluriformes, as well as in several other families of this order (OLIVEIRA and GOSZTONYI 2000). Additionally, several species of the Trichomycteridae family, a primitive sister group of Callichthyidae (DE PINNA 1998) have 2n=54 chromosomes, single Ag-NORs and small C-band positive segments in centromeric position (LIMA and GALETTI JR. 1990; Torres et al. 1998; Borin and Santos 1999; Gonzo et al. 2000). Thus, the cytogenetic characteristics found in C. callichthys, with diploid number ranging from 2n=52 and 2n=58, being three samples with 2n=56 (SANCHEZ and FENоссню 1996; present study), several M/SM chromosomes, a single chromosome pair with Ag-NORs, and small C-band positive segments located in centromeric position, permit to suggest that *Callichthys* may be the most primitive genus of the Callichthyinae family. This hypothesis is in accordance with the phylogenetic position of this genus in a morphological phylogeny constructed by Reis (1998).

The cytogenetic characteristics of *Dianema*, as 2n=60-62 chromosomes, many ST/A chromosomes, a single chromosome pair with Ag-NORs, the pattern of distribution of C-band positive segments, and its low amount of nuclear DNA (OL-IVEIRA et al. 1993) are similar to those observed in Hoplosternum littorale and Hoplosternum sp. (OLIVEIRA et al. 1993), suggesting a close relationship between these genera, as proposed by OL-IVEIRA et al. (1993). These data are in accordance with the phylogeny proposed by REIS (1998), in which these two genera belong to a monophyletic group. Additionally, studies with the amount of nuclear DNA content showed that Dianema urostriata had the smallest value of the subfamily, 1.18 ± 0.07 pg of DNA by diploid nuclei, followed by *Hoplosternum* sp., 1.36 ± 0.11 , and two samples of C. callichthys, 1.89 ± 0.24 and 1.94 ± 0.15 (OL-IVEIRA et al. 1993), reinforcing the hypothesis that Dianema and Hoplosternum are more related between themselves than to Callichthys (Reis 1998).

The cytogenetic analyses conducted in the Callichthyinae subfamily showed the occurrence of two groups of species. The first group is composed by the genus Callichthys, that ranges from 2n=52 to 2n=58 chromosomes and many M/SM chromosomes. The second group include the other genera of the subfamily, being characterized by diploid number that ranges from 2n=60 to 2n=66 chromosomes and high number of A chromosomes. Contrarily to the extensive chromosome rearrangements observed in the Corydoradinae subfamily, in the Callichthyinae subfamily a small number of chromosome changes seem to have occurred in the evolutionary history of the group. On the other hand, the karyotypic variability observed among the Callichthyinae suggest the existence of several undescribed species in this subfamily.

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