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Cytogenetic analysis in *Tetragonopterus franciscoensis* (Characiformes): another piece to the karyoevolutionary puzzle of tetra fishes

MAURICIO BARROS FERNANDES, JAMILLE DE ARAÚJO BITENCOURT, JOANDSON CALIXTO DOS SANTOS, JOSÉ HENRIQUE GALDINO*, PAULO ROBERTO ANTUNES DE MELLO AFFONSO

Department of Biological Sciences, Universidade Estadual do Sudoeste da Bahia (UESB), Jequié - BA, Brazil

*Corresponding author. E-mail: galdino.jhsbg@gmail.com

Abstract. *Tetragonopterus* is a taxonomically complex genus in Characidae, being currently represented by nine species according to integrative approaches. One of them, *T. franciscoensis* was recently validated in rivers from northeastern Brazil. Even though molecular and morphological data have been collected in *Tetragonopterus*, the cytogenetic analyses in this group are scarce despite of the role of chromosomal variation in speciation. Herein, we present the first detailed karyotypic study in *T. franciscoensis* along with a comparative analysis with published cytogenetic data in characin fish. All specimens shared $2n=52$ distributed in 12 metacentric (m), 12 submetacentric (sm), and 28 subtelo/acrocentric (st/a) chromosomes for both sexes as well as single nucleolus organizer regions on short arms of pair 8 and several GC-rich sites. The mapping of telomeric sequences (TTAGGG)_n revealed no telomeric interstitial signals. While subtle cytogenetic differences were observed between samples from northeastern basins in Brazil, corroborating a recent genetic divergence, distinct karyotypes were detected in relation to congeneric taxa from other Brazilian regions. Therefore, the origin of large biarmed pairs in species with low $2n$ values should be related to occurrence of centric fusions.

Keywords: Characidae, Characins, cytotaxonomy, neotropical fish, Tetragonopterinae.

INTRODUCTION

The genus *Tetragonopterus* (Characidae) was proposed by Cuvier (1816) to describe the species *T. argenteus* based on a unique specimen from South America. In the second half of the 19th century, Günther (1864) added 32 new species to this taxon and proposed the subfamily Tetragonopterinae which would include most of small characins or tetras (e.g., *Astyanax*, *Hemigrammus*, *Moenkhausia*, *Psalidodon*).

Over the following decades, the group was extensively revised and it turned to be one of the most intriguing taxa among Characidae. In a series of studies carried out by Carl H. Eigenmann, several species previously allo-

cated in *Tetragonopterus* were reassigned to different genera, like *Bryconamericus*, *Ctenobrycon*, and *Deuterodon* (Eigenmann, 1917; Eigenmann, 1918; Eigenmann, 1921; Eigenmann and Myers, 1929). Later, the number of species in *Tetragonopterus* was reduced to four evolutionary units, comprising *T. argenteus*, *T. chalceus*, *T. gibbosus*, and *T. huberi*. On that occasion, the reassignment of *T. georgiae* and *T. rarus* to *Moenkhausia*, for example, was justified by the lack of a complete lateral line greatly bent downwards at the anterior portion, a common feature of *Tetragonopterus*. Follow-up taxonomic reviews reallocated *T. argenteus* and *T. chalceus* as the only representatives of this genus (Reis *et al.*, 2003). However, this scenario has changed considerably, as DNA-based studies provided important insights about the taxonomic relationships of *Tetragonopterus* and other tetras (Araújo and Lucinda, 2014; Mirande, 2019).

Accordingly, molecular analysis recognized eight previously described species in *Tetragonopterus* (*T. anostomus*, *T. araguaiensis*, *T. argenteus*, *T. carvalhoi*, *T. chalceus*, *T. denticulatus*, *T. georgiae*, and *T. rarus*) and cases of cryptic diversity (Silva *et al.*, 2016). These authors revealed that the populations of *T. chalceus* from São Francisco, Paraguaçu, and Itapicuru river basins actually encompassed a distinct species, referred to as *T. franciscoensis* (Silva *et al.*, 2016). In addition, three new species were also described (*T. jurema*, *T. kulene*, and *T. ommotus*) and new evidence reallocated *Moenkhausia georgiae* back to *Tetragonopterus* (*T. georgiae*), as also supported by other authors (Silva *et al.*, 2016; Melo *et al.*, 2016; Terán *et al.*, 2020).

Even though the abovementioned studies were particularly informative to resolve the taxonomic uncertainties in *Tetragonopterus*, cytogenetic analyses that could add new pieces of evidence to this subject remain limited to a few reports based on conventional analyses in *T. argenteus* Cuvier, 1816 and *T. chalceus* Spix & Agassiz, 1829. Both species shared a modal diploid number of $2n = 52$, a single NOR system and few heterochromatin regions, but they diverge in their karyotype formulae (Portela *et al.*, 1988; Alberdi and Fenocchio, 1997). Interestingly, populations of *T. argenteus* from Cuiabá River were differentiated by the presence of two cytotypes (1 and 2). While the cytotype 1 is represented by specimens with $2n=50$ and a karyotype of $14m+4sm+4st+28a$, the cytotype 2 presents $2n=52$ distributed into $14m+4sm+4st+30a$ chromosomes (Miyazawa, 2015).

A striking cytogenetic feature commonly reported in small characins is the presence of a large first metacentric pair when compared to other chromosomes in the karyotype (Scheel, 1973). In fact, this metacentric

pair and a modal number of $2n=50$ have been regarded as plesiomorphies for this fish group (Morelli *et al.*, 1983; Portela-Castro *et al.*, 1998; Tenório *et al.*, 2013), being also observed in Bryconidae (Almeida-Toledo *et al.*, 1996; Mariguela *et al.*, 2010; Yano *et al.*, 2021).

In turn, the highly conserved morphology of small characins, including *Tetragonopterus* (Eigenmann, 1917), indicates that species complexes or cryptic species might be present, thus hindering reliable estimates of richness and endemism rates in these Neotropical fishes. In this context, cytogenetic methods can help reveal such overlooked diversity, as exemplified by studies in the genus *Psalidodon* (e.g. Bertaco *et al.*, 2006; Ferreira-Neto *et al.*, 2012). Therefore, the goal of the present study was to report the first detailed cytogenetic characterization of *T. franciscoensis* from an isolated drainage from Northeastern Brazil to shed some light on the taxonomy and species delimitation in *Tetragonopterus*. In addition, we carried out a comprehensive comparative cytogenetic analysis in characin species to provide insights about the karyoevolutionary trends in the subfamily Tetragonopterinae.

MATERIAL AND METHODS

Thirteen individuals of *T. franciscoensis* Silva, Melo, de Oliveira & Benine, 2016 (8 males and 5 females) were collected at the Itapicuru-Mirim River (Itapicuru River Basin) in the municipality of Tucano, state of Bahia, northeastern Brazil ($11^{\circ}12'15.3''S/40^{\circ}29'15.1''W$) (Fig. 1). The collection license was granted by the Instituto Chico Mendes de Conservação da Biodiversidade (ICMBio/SISBIO n. 26752-2). The procedures and experiments were approved by the Committee of Ethics in Experimentation with Animals from the State University of Southwestern Bahia (CEUA/UESB 32/2013).

To stimulate cell division, the fish specimens were inoculated with fungal antigens and kept in tanks for 48 to 72 hours (Lee and Elder, 1980). Afterwards, the specimens were euthanized in cold water (Blessing *et al.*, 2010), and the anterior kidney was removed to obtain metaphase cells, according to Netto *et al.*, (2007). The cell suspension containing the mitotic chromosomes were dropped on glass slides, air dried and stained with 10% Giemsa in phosphate buffer (pH 6.8).

The heterochromatin was visualized by C-banding technique (Sumner, 1972), and active nucleolar organizer regions (Ag-NOR) were detected by silver staining (Howell and Black, 1980). Sequential staining with the base-specific fluorochromes Chromomycin A₃ (CMA₃) and 4'-6-diamino-2-fenilindole (DAPI) to detect GC-

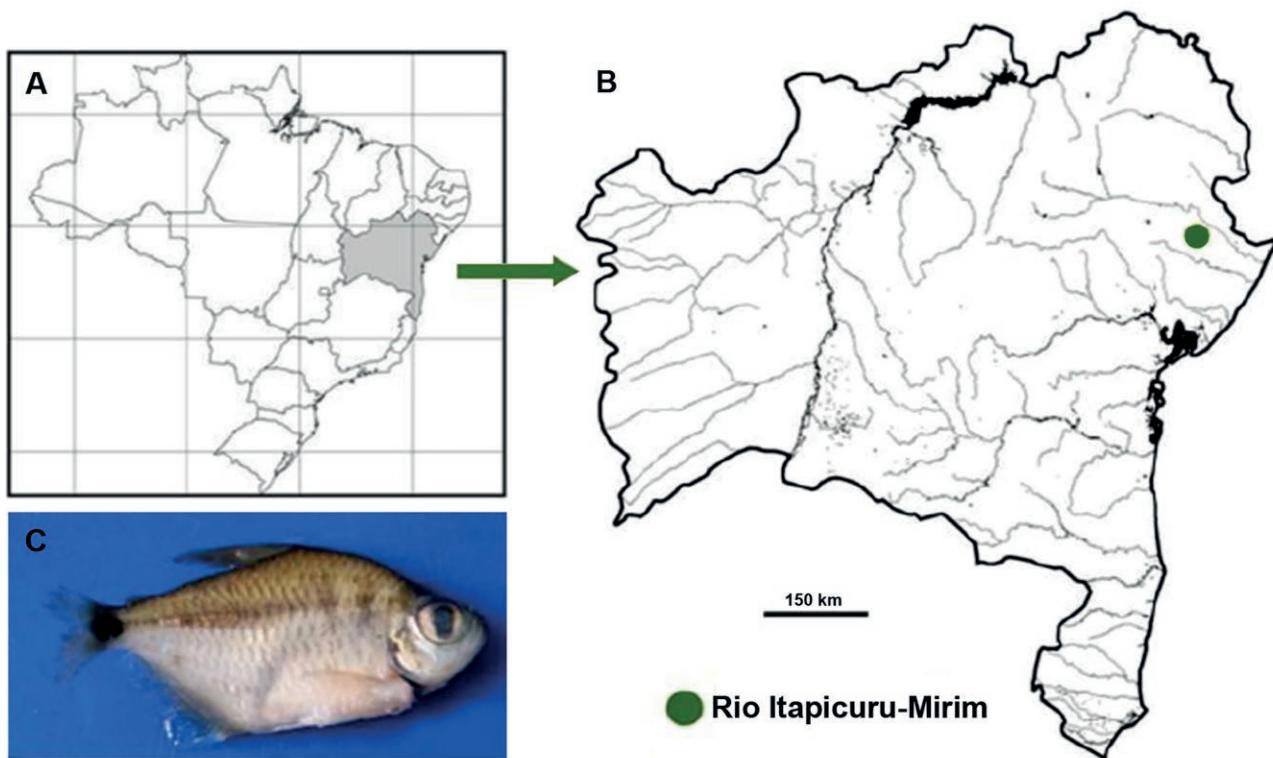


Figure 1. Map of Brazil highlighting the state of Bahia (a) and the collection site in Itapicuru-Mirim River (b) of *T. franciscoensis* (c).

and AT-rich regions, respectively, were carried out according to Schmid (1980).

The physical mapping of telomeres was performed based on fluorescence *in situ* hybridization (FISH) according to Pinkel *et al.* (1986) under high stringency (77%) conditions to evaluate the putative presence of internal telomere sequences (ITS) that could reveal structural rearrangements. The telomere (TTAGGG)_n probes were obtained via PCR without template DNA (Ijdo *et al.*, 1991). The probes were labeled with digoxigenin-11-dUTP and detected with anti-digoxigenin-Rhodamine conjugate, according to the manufacturer's instructions (Roche). The chromosomes were counterstained with DAPI and the slides were mounted in a Vectashield medium.

A mean number of 10 metaphase spreads per specimen were analyzed using an epifluorescence microscope (Olympus BX-51) attached to a digital camera and equipped with the software Image-Pro Plus® v. 6.2 for photo documentation. The chromosomes were measured using the software Easy Idio 1.0 (Diniz and Melo, 2006). Then, they were classified according to their arm ratio (Levan *et al.* 1964), and the chromosomal pairs were systematically organized into karyotypes in decreasing size order within each morphological category.

RESULTS

A modal diploid number of $2n = 52$ was observed in all specimens of *T. franciscoensis*, while the karyotype was invariably organized into 12 metacentric (m), 12 submetacentric (sm), and 28 subtelocentric/acrocentric (st/a) chromosomes (Figure 2a). No heteromorphic sex chromosomes were detected.

The silver staining revealed a single NOR-bearing pair (8) with heteromorphic ribosomal cistrons at interstitial regions on short arms. On the other hand, the C-banding revealed few heterochromatin blocks restricted to centromeres (Figure 2b). The GC-rich sites (CMA_3^+ /DAPI) were coincident with Ag-NORs on pair 8 (Figure 3). Furthermore, additional CMA_3 signals were observed in, at least, three other chromosomal pairs (Figure 3). The mapping of (TTAGGG)_n sequences by FISH revealed conspicuous signals on telomeres of all chromosomes and no internal telomere sequences (ITS) (Figure 4).

DISCUSSION

The karyotype macrostructure of *T. franciscoensis* ($2n=52$ and a karyotype formula of $12m+12sm+28st/a$)

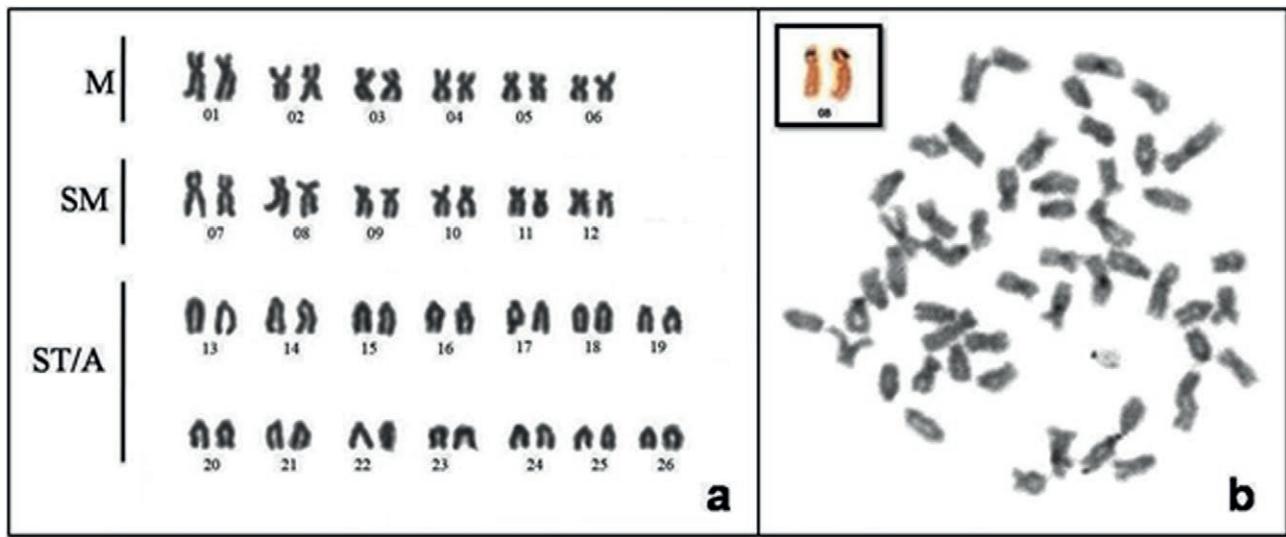


Figure 2. Giemsa-stained (a) and C-banded (b) karyotypes of *T. franciscoensis*. The NOR-bearing pair after silver nitrate is shown in box.

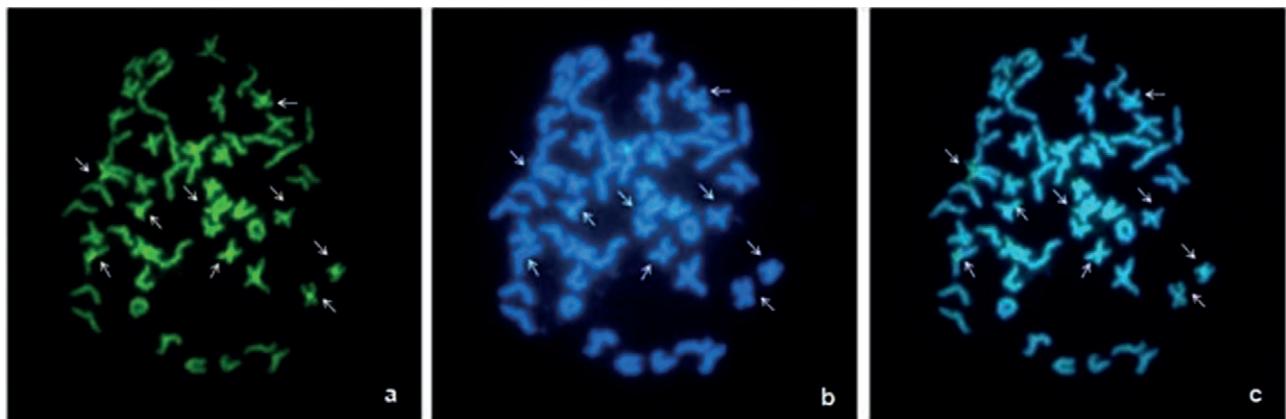


Figure 3. Somatic metaphases after CMA₃ (a); DAPI (b); and overlapped CMA₃/DAPI (c) staining, showing the GC-rich (CMA₃⁺/DAPI⁻) signals indicated by arrows.

is similar to that reported in populations of *T. chalceus* (=*T. franciscoensis* sensu Silva *et al.*, 2016) from São Francisco River (26m/sm+26st/a) (Portela *et al.*, 1988). The only difference refers to the presence of an additional subtelocentric/acrocentric pair in specimens from Itapicuru-Mirim (present study). This result suggests a genetic divergence among these lineages from each hydrographic system driven by pericentric inversions in a chromosome pair. Nevertheless, artifactual effects could also account for these such as distinct levels of chromosome condensation or the criteria for determining the chromosomal morphology between authors.

On the other hand, remarkable macrostructural differences are observed when the cytogenetic data in *T. franciscoensis* from the present study are compared

to those reported in closely related species, such as *T. argenteus* from Paraná (16m/sm+2st+34a), Paraguay (14m+4sm+4st+28a), and (De La Plata) river basins (Alberdi and Fenocchio, 1997; Miyazawa, 2015). In the latter, the specimens presented interindividual variation in both number and morphology of chromosomes (2n=50, 14m+4sm+4st+28a and 2n=52, 16m+4sm+4st+28a) (Miyazawa, 2015; Supplementary Table 1). These findings indicate that inversions and fusions/fissions are involved in the karyoevolution of *Tetragonopterus* and that cryptic species are likely to be present in this group, as commonly observed in small characins (Medrado *et al.*, 2018).

Furthermore, *T. franciscoensis* lacks the typical large metacentric pair described in other tetras, a condition putatively associated with the presence of 2n=52

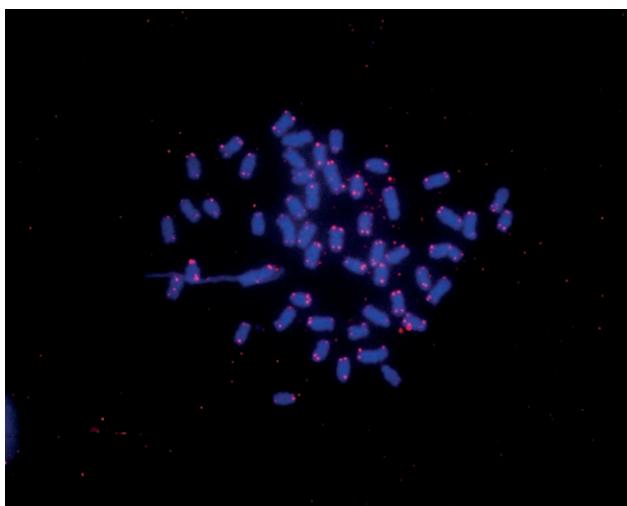


Figure 4. Metaphase of *T. franciscoensis* after FISH with (TTAGGG)_n probes, revealing the positive signals (in magenta) on telomeres.

(Figure 2a), thus diverging from the pattern observed in several genera of Characidae like *Astyanax*, *Psalidodon*, *Moenkhausia*, and *Cheirodon* (Tenório *et al.*, 2013; Soto *et al.*, 2018; Nascimento *et al.*, 2020). Such difference reinforces the divergence of *T. franciscoensis* and congeners in relation to other small characin lineages, corroborating their allocation in a distinct subfamily (Tetragonopterinae) (Mirande, 2018; Terán *et al.*, 2020).

In fact, a distinctive first large metacentric pair is also found in representatives from other closely related and basal families of Characiformes (Supplementary Table 1), such as Bryconidae (Almeida-Toledo *et al.*, 1996, Mariguela *et al.*, 2010; Silva *et al.*, 2012), indicating that this is a plesiomorphic condition. Moreover, this condition (presence or absence of large metacentric pairs) varies remarkably among distinct taxonomic units in Characidae. Such variation has been reported even within some genera such as *Astyanax*, *Psalidodon*, and *Hyphessobrycon*, and within species, like *Bryconamericus* aff. *exodon* and *Bryconamericus* aff. *iheringii*, indicating putative species complexes or cryptic diversity (Supplementary Table 1).

On the other hand, the absence of a long metacentric pair appears to be ubiquitous in *Odontostilbe*, *Piabina*, *Serrapinnus*, and *Knodus* (Supplementary Table 1). Moreover, according to the present revision, the lack of this large metacentric pair is correlated with species characterized by 2n=52 (Supplementary Table 1). Therefore, it is reasonable to hypothesize that independent chromosomal fusion events could account for the very large size of the first pair of biarmed chromosomes and the reduction of diploid numbers (2n < 50) in char-

acins. However, these findings are insufficient to fully understand the karyoevolutionary trends in Characidae because several genera and species in this family remain poorly studied in relation to their cytogenetic traits. Therefore, further basic chromosomal studies should be carried out to test the role of centric fusions in the karyoevolution of small characins and the utility of the largest metacentric pair as a cytotaxonomic marker in tetras.

Similarly, the number and distribution of NORs in *T. franciscoensis* (Figure 2b) resembles that of *T. chalceus* (Portela *et al.*, 1988) and *T. argenteus* (Miyazawa, 2015), following a common trend among characins (Medrado *et al.*, 2008). In addition, the presence of GC-rich (CMA₃⁺) sites co-located with NORs are considered a basal trait for fish and amphibianli (Schmid, 1980; Tenório *et al.*, 2013; Monteiro *et al.*, 2022). On the other hand, the presence of additional GC-rich sites at centromeric regions (Figure 3) represent a unique and putatively apomorphic condition since AT-rich sites near centromeres are more frequently reported in small characins (Sánchez *et al.*, 2021), thus indicating a heterogeneous composition of satellite DNAs. These results show the importance of detailed chromosomal analyses to infer the dynamics of genome organization and the role of microarrangements in speciation of tetra fishes.

The mapping of telomeric sequences on chromosomes of *T. franciscoensis* (Figure 4) followed the expected pattern in vertebrates, revealing positive signals at terminal portions of chromosomal pairs (Meyne *et al.*, 1989; Ferro *et al.*, 2003; Schmid *et al.*, 2006) and no evidence of ITS. Nonetheless, this pattern should not reject the occurrence of chromosomal rearrangements in the analyzed species. Actually, ITS are often lost or degenerated in rearranged chromosomes, particularly when the chromosomal changes have occurred in early stages of differentiation among clades (Meyne *et al.*, 1990; Bolzan, 2017).

In general, the present study revealed subtle cytogenetic differences in *Tetragonopterus* from São Francisco and Itapicuru River basins in northeastern Brazil, contrasting with the distinct karyotypes of congeneric species from other Brazilian regions (e.g., *T. argenteus*). These findings provide additional support to the validation of these populations as *T. franciscoensis* as proposed by morphological data (Silva *et al.*, 2016). At last, the lack of the typical large metacentric pair and the predominance of 2n=52 in *Tetragonopterus* when compared to other small characins reinforced the status of Tetragonopterinae as a monophyletic subfamily. In addition, cytotaxonomic markers were reported for *T. franciscoensis* that can be properly used to resolve taxonomic uncertainties in Neotropical tetras.

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REFERENCES

- Alberdi AJ, Fenocchio, AS. 1997. Karyotypes of Five Tetragonopterinae Species (Pisces, Characidae) from Argentina. *Cytologia*. 62:171-176.
- Almeida-Toledo LFD, Bigoni AP, Bernardino G, Foresti F, Toledo-Filho SDA. 1996. Karyotype and NOR conservatism with heterochromatin reorganization in Neotropical Bryconids. *Caryologia*. 49(1):35-43.
- Alves AL, Martins-Santos IC. 2002. Cytogenetics studies in two populations of *Astyanax scabripinnis* with $2n=48$ chromosomes (Teleostei, Characidae). *Cytologia*. 67(2):117-122.
- Araujo L, Lucinda PHF. 2014. A new species of the genus *Tetragonopterus* Cuvier, 1816 (Ostariophysi: Characiformes: Characidae) from the rio Tocantins drainage, Brazil. *Neotropical Ichthyology*. 12:309-315.
- Arefjev VA. 1990. Problems of karyotypic variability in the family Characidae (Pisces, Characiformes) with the description of somatic karyotypes for six species of tetras. *Caryologia*. 43 (3-4):305-319.
- Barreto SB, Cioffi MB, Medrado AS, Silva AT, Affonso, PR, Diniz D. 2016. Allopatric chromosomal variation in *Nematocharax venustus* Weitzman, Menezes & Britski, 1986 (Actinopterygii: Characiformes) based on mapping of repetitive sequences. *Neotropical Ichthyology*. 14(2):e150141.
- Blessing JJ, Marshall JC, Balcombe SR. 2010. Humane killing of fishes for scientific research: a comparison of two methods. *Journal of Fish Biology*. 76(10):2571-2577.
- Bolzan DA. 2017. Interstitial telomeric sequences in vertebrate chromosomes: Origin, function, instability and evolution. *Mutation Research/Reviews in Mutation Research*. 773:51-65.
- Capistano TG, Portela-Castro ALDB, Júlio-Junior HF. 2008. Chromosome divergence and NOR polymorphism in *Bryconamericus* aff. *iheringii* (Teleostei, Characidae) in the hydrographic systems of the Paranaapanema and Ivaí Rivers, Paraná, Brazil. *Genetics and Molecular Biology*. 31:203-207.
- Carvalho ML, Oliveira C, Foresti F. 2002. Cytogenetic analysis of five species of the subfamily Tetragonopterinae (Teleostei, Characiformes, Characidae). *Caryologia*. 55(3):181-188.
- Centofante L, Bertollo LAC, Miyazawa CS, Moreira-Filho, O. 2003. Chromosomal differentiation among allopatric populations of *Hyphessobrycon anisitsi* (Pisces, Tetragonopterinae). *Cytologia*. 68(3):283-288.
- Coutinho-Sanches N, Dergam JA. 2015. Cytogenetic and molecular data suggest *Deuterodon pedri* Eigenmann, 1907 (Teleostei: Characidae) is a member of an ancient coastal group. *Zebrafish*. 12(5):357-365.
- Cunha MS. 2021. Repetitive DNA mapping on *Oligosarcus acutirostris* (Teleostei, Characidae) from the Paráiba do Sul River Basin in southeastern Brazil. *Caryologia*. 74(4):121-128.
- Cuvier G. 1816. Le règne animal distribué d'après son organisation pour servir de base à l'histoire naturelle des animaux et d'introduction à l'anatomie comparée. Les reptiles, les poissons, les mollusques et les annélides. Edition 1 v.2., Déterville, Paris, xviii+532.
- Da Silva LLL, Dos Santos AR, Giuliano-Caetano L, Dias, AL. 2016. Chromosomal characterization in two species of an *Astyanax bimaculatus* complex (Characidae, Characiformes) using different techniques of chromosome banding. *Cytotechnology*. 68:1277-1286.
- Da Silva LLL, Giuliano-Caetano L, Dias AL. 2014. Karyotypic diversity in a population of *Bryconamericus* aff. *iheringii* (Characidae). *Genetics and Molecular Research*. 13(1):2069-2081.
- Da Silva LLL, Giuliano-Caetano L, Dias AL. 2012. Chromosome studies of *Astyanax jacuhiensis* Cope, 1894 (Characidae) from the Tramandai River Basin, Brazil, using *in situ* hybridization with the 18S rDNA probe, DAPI and CMA3 staining. *Folia Biologica (Kraków)*. 60(3-4):135-140.
- Da Silva LL, Dos Santos AR, Giuliano-Caetano L, Dias AL. 2016. Chromosomal characterization in two species of an *Astyanax bimaculatus* complex (Characidae, Characiformes) using different techniques of chromosome banding. *Cytotechnology*. 68:1277-1286.
- Diniz D, Xavier PM. 2006. Easy Idio. Available: <http://geocities.yahoo.com.br/easyidio/>.
- Domingues MDS, Vicari MR, Abilhoa V, Wamser JP, Cestari MM, Bertollo LA, Artoni RF. 2007. Cytogenetic and comparative morphology of two allopatric populations of *Astyanax altiparanae* Garutti & Britski, 2000 (Teleostei: Characidae) from upper rio Paraná basin. *Neotropical Ichthyology*. 5:37-44.

- Dos Santos R, Medrado AAS, Diniz D, Oliveira C, Affonso PRAM. 2016. ZZ/ZW sex chromosome system in the endangered fish *Lignobrycon myersi* Miranda-Ribeiro, 1956 (Teleostei, Characiformes, Triportheidae). Comparative Cytogenetics. 10(2):245.
- Dos Santos AR, Rubert M, Giuliano-Caetano L, Dias, AL. 2012. Sympatric occurrence of four cytotypes and one extra chromosome in *Bryconamericus ecai* (Characidae): 18S rDNA polymorphism and heterochromatin composition. *Hereditas*. 149(1):24-33.
- Duarte MA, Giugliano LG, de Aquino PDP, Grisolia CK, Milhomem-Paixão SSR. 2018. Cytogenetic studies in *Hasemania crenuchoides* (Characiformes: Characidae) and molecular investigations into kinship relationships of the genus. *Caryologia*. 71(4):446-452.
- Eigenmann CH. Catalogue of the fresh-water fishes of tropical and south temperate America. 1910.
- Eigenmann CH. 1903. New genera of South American fresh-water fishes, and new names for some old genera. Smithsonian Miscellaneous Collections.
- Eigenmann CH. 1908. Preliminary descriptions of new genera and species of tetragonopterid characins (Zoölogical Results of the Thayer Brazilian expedition.). Bulletin of the Museum of Comparative Zoology. 52(6):91-106.
- Eigenmann CH. 1917. The American Characidae, Part 1. Memoirs of the Museum of Comparative Zoology. 43:1-102.
- Eigenmann CH. 1918. The American Characidae, Part 2. Memoirs of the Museum of Comparative Zoology. 43:103-208.
- Eigenmann CH. 1921. The American Characidae, Part 3. Memoirs of the Museum of Comparative Zoology. 43:209-310.
- Eigenmann CH, Myers G. 1929. The American Characidae, Part 5. Memoirs of the Museum of Comparative Zoology. 43:429-558.
- Fernandes CA, Piscor D, Bailly D, Da Silva VFB, Martins-Santos IC. 2010. Cytogenetic studies comparing three Characidae fish species from the Iguaçemi River Basin, Brazil. *Cytologia*. 75(4):329-333.
- Ferreira Neto M, Vicari, MR, Camargo EFD, Artoni RF, Moreira-Filho O. 2009. Comparative cytogenetics among populations of *Astyanax altiparanae* (Characiformes, Characidae, Incertae sedis). *Genetics and Molecular Biology*. 32:792-796.
- Ferro MAD, Moreira-Filho O, Bertollo CAL. 2003. B Chromosome Polymorphism in the Fish, *Astyanax scabripinnis*. *Genetica*. 119:147-53.
- Foresti F, Almeida-Toledo LF, Toledo SA. 1989. Supernumerary chromosome system, C-banding pattern characterization and multiple nucleolus organizer regions in *Moenkhausia sanctaefilomenae* (Pisces, Characidae). *Genetica*. 79(2):07-114.
- Freitas PD, Galetti-Jr PM. 1998. Chromosome characterization of a neotropical fish *Poptella paraguayensis* from Paraguay river basin (Stethaprioninae, Characidae). *Cytologia*. 63(1):73-77.
- Gavazzoni M, Paiz LM, Oliveira CA, Pavanelli CS, Graca WJ, Margarido, VP. 2018. Morphologically cryptic species of the *Astyanax bimaculatus* "caudal peduncle spot" subgroup diagnosed through cytogenetic characters. *Zebrafish*. 15(4):382-388.
- Goes CAG, Daniel SN, Piva LH, Yasui GS, Artoni RF, Hashimoto DT, Porto-Foresti F. 2020. Cytogenetic markers as a tool for characterization of hybrids of *Astyanax* Baird & Girard, 1854 and *Hyphessobrycon* Eigenmann, 1907. Comparative Cytogenetics. 14(2):231.
- Günther ACLG. 1864. Catalogue of the fishes in the British Museum. Catalogue of the Physostomi, containing the families Siluridae, Characinidae, Haplochitonidae, Sternopychidae, Scopelidae, Stomiataidae in the collection of the British Museum. Catalogue of the fishes in the British Museum. 5:1-455.
- Guimarães IN, Foresti de Almeida-Toledo L, Oliveira C, Foresti F, De Almeida Toledo Filho S. 1995. Cytogenetic studies of three species of Glandulocaudinae (Pisces, Characiformes, Characidae). *Revista Brasileira de Genética*. 18(2):185-189.
- Hattori RS, Zambelli FDSM. 2007. Foresti, A.T.L. Karyotype characterization and gene mapping of 5S and 18S rDNA in three species of *Oligosarcus* (Teleostei: Characidae). *Caryologia*. 60(4):372-378.
- Howell WMT, Black DA. 1980. Controlled silver-staining of nucleolus organizer regions with a protective colloidal developer: a 1-step method. *Experientia* 36(8):1014-1015.
- Ijdo JW, Wells RA, Baldini A, Reeders ST. 1991. Improved telomere detection using a telomere repeat probe (TTAGGG) n generated by PCR. *Nucleic acids research*. 19(17):4780.
- Kantek DLZ, Noleto RB, Maurutto FAM, Bertollo LAC, Moreira-Filho O, Cestari MM. 2008. Cytotaxonomy of *Astyanax* (Characiformes, Characidae) from the Upper Iguaçu River Basin: confirmation of the occurrence of distinct evolutionary units. *Journal of Fish Biology*. 73(8):2012-2020.
- Kavalco KF, Pazza R, Almeida-Toledo LF. 2008. *Astyanax bockmanni* Vari and Castro, 2007: an ambiguous karyotype in the *Astyanax* genus. *Genetica*. 136:135-139.
- Kavalco KF, Pazza R, Bertollo LA, Moreira-Filho O. 2005. Molecular cytogenetics of *Oligosarcus hepsetus* (Teleostei, Characiformes) from two Brazilian locations. *Genetica*. 124:85-91.

- Kavalco KE, Moreira-Filho O. 2003. Cytogenetical analyses in four species of the genus *Astyanax* (Pisces, Characidae) from Paraíba do Sul River basin. *Caryologia*. 56(4):453-461.
- Klassmann JP, Dos Santos ICM. 2017. Karyotype diversity in *Astyanax scabripinnis* (Pisces, Characiformes) from the São Francisco River basin, Brazil. *Cytologia*. 82(2):199-203.
- Krinski D, Centofante L, Miyazawa CS. 2008. Multiple NORs in *Knodus* cf. *chapadae* (Pisces: Characidae: *incertae sedis*) from Upper Paraguay Basin, Mato Grosso State, Brazil. *The Nucleus*. 51(3):229-238.
- Krinski D, Miyazawa CS. 2012. First description of the karyotype and Ag-NORs localization of *Brachychalcinus retrospina* (Pisces: Characidae: Stethaprioninae) from Upper Paraguay Basin, Mato Grosso State, Brazil. *Caryologia*. 65(1):82-85.
- Lee MR, Elder FFB. 1980. Yeast stimulation of bone marrow mitosis for cytogenetic investigations. *Cytogenetic and Genome Research*. 26(1):36-40.
- Levan, A. 1964. Nomenclature for centromeric position on chromosomes. *Hereditas*. 52:201-220.
- Maistro EL, Foresti F, Oliveira C, de Almeida Toledo LF. 1992. Occurrence of macro B chromosomes in *Astyanax scabripinnis paranae* (Pisces, Characiformes, Characidae). *Genetica*. 87:101-106.
- Mari-Ribeiro IP, Scorsim B, Oliveira AVD, Portela-Castro ALDB. 2022. Cytogenetic and Molecular Characterization of *Oligosarcus pintoi* (Characidae): A New Record of Supernumerary Chromosome in this Species. *Zebrafish*. 19(2):71-80.
- Mariguela T, Nirchio M, Ron E, Gaviria J, Foresti F, Oliveira C. 2010. Cytogenetic characterization of *Brycon amazonicus* (Spix et Agassiz, 1829) (Teleostei: Characidae) from Caicara del Orinoco, Venezuela. *Comparative Cytogenetics*. 4(2):185-193.
- Martinez ER, Alves AL, Silveira SM, Foresti F, Oliveira C. 2012. Cytogenetic analysis in the *incertae sedis* species *Astyanax altiparanae* Garutti and Britzki, 2000 and *Hypseobrycon eques* Steindachner, 1882 (Characiformes, Characidae) from the upper Paraná River basin. *Comparative Cytogenetics*. 6(1):41.
- Medrado AS, Affonso PRMA, Carneiro PL, Vicari MR, Artoni RF, Costa MA. 2015. Allopatric divergence in *Astyanax* aff. *fasciatus* Cuvier, 1819 (Characidae, Incertae sedis) inferred from DNA mapping and chromosomes. *Zoologischer Anzeiger-A Journal of Comparative Zoology*. 257:119-129.
- Medrado AS, Figueiredo AVA, Waldschmidt AM, Affonso PRAM, Carneiro PLS. 2008. Cytogenetic and morphological diversity in populations of *Astyanax fasciatus* (Teleostei, Characidae) from Brazilian north-eastern river basins. *Genetics and Molecular Biology*. 31:208-214.
- Melo BF, Benine RC, Silva GS, Avelino GS, Oliveira C. 2016. Molecular phylogeny of the Neotropical fish genus *Tetragonopterus* (Teleostei: Characiformes: Characidae). *Molecular Phylogenetics and Evolution*. 94:709-717.
- Mestriner CA, Galetti PM, Valentini SR, Ruiz IR, Abel LD, Moreira-Filho O, Camacho JP. 2000. Structural and functional evidence that a B chromosome in the characid fish *Astyanax scabripinnis* is an isochromosome. *Heredity*. 85(1):1-9.
- Mendes MM, Rosa R, Giuliano-Caetano L, Dias AL. 2011. Karyotype diversity of four species of the *incertae sedis* group (Characidae) from different hydrographic basins: analysis of AgNORs, CMA3 and 18S rDNA. *Genetics and Molecular Research*. 10(4):3596-3608.
- Meyne J, Baker RJ, Hobart HH, Hsu TC, Ryder OA, Ward OG, Moyzis RK. 1990. Distribution of non-telomeric sites of the (TTAGGG) n telomeric sequence in vertebrate chromosomes. *Chromosoma*. 99:3-10.
- Meyne J, Ratliff RL, MoYzIs RK. 1989. Conservation of the human telomere sequence (TTAGGG) n among vertebrates. *Proceedings of the National Academy of Sciences*. 86(18):7049-7053.
- Mirande JM. 2019. Morphology, molecules and the phylogeny of Characidae (Teleostei, Characiformes). *Cladistics*. 35(3):282-300.
- Miyazawa C. 2015. Diferentes citótipos de *Tetragonopterus argenteus* Cuvier, 1817 (Characiformes, Tetragonopterinae) e descrição cariotípica de *Markiana nigripinnis* Perugia, 1891 (Characiformes, Characidae, *incertae sedis*) de rios da bacia do rio Paraguai. *Scientia Vitae*. 2:17- 27.
- Monteiro ABGF, Takagui FH, Baldissera JNDC, Jerep FC, Giuliano-Caetano L. 2022. Classical and molecular cytogenetics of *Markiana nigripinnis* (Pisces-Characiformes) from Brazilian Pantanal: a comparative analysis with cytotaxonomic contributions. *Biologia*. 77(8):2371-2382.
- Moreira PWA, Bertollo LAC, Moreira-filho O. 2007. Comparative cytogenetics between three Characidae fish species from the São Francisco River basin. *Caryologia*. 60(1-2):64-68.
- Morelli S, Bertollo LAC, Foresti F, Moreira-Filho O, De Almeida Toledo FS. 1983. Cytogenetic considerations on the genus *Astyanax* (Pisces, Characidae). I. Karyotypic variability. *Caryologia*. 36(3):235-244.
- Nanda I, Schneider-Rasp S, Winking H, Schmid M. 1995. Loss of telomeric sites in the chromosomes of *Mus musculus domesticus* (Rodentia: Muridae) dur-

- ing Robertsonian rearrangements. Chromosome Research. 3:399-409.
- Nascimento CND, Troy WP, Alves JCP, Carvalho ML, Oliveira C, Foresti F. 2020. Molecular cytogenetic analyses reveal extensive chromosomal rearrangements and novel B chromosomes in *Moenkhausia* (Teleostei, Characidae). Genetics and Molecular Biology. 43(4): e20200027.
- Netto MRCB, Paula E, Affonso PRAMA. 2007. standard protocol for obtaining fish chromosomes under post-mortem conditions. Micron. 38(3):214-217.
- Nishiyama PB, Rossi MM, Porto FE, Borin LA, Castro ALDBP, dos Santos ICM. 2015. Estudos citogenéticos em espécies de peixes de riachos: *Hypessobrycon vinaceus*, *Bryconamericus* aff. *iheringii* e *Odontostilbe pequira* (PISCES: CHARACIDAE). Evolução e Conservação da Biodiversidade. 6(1):13-22.
- Pacheco RB, Giuliano-Caetano L, Julio Junior HF, Dias AL. 2010. Cytogenetic data on *Astyanax jacuhiensis* (Characidae) in the lago Guaíba and tributaries, Brazil. Neotropical Ichthyology. 8:667-671.
- Paintner-Marques TR, Giuliano-Caetano L, Dias AL. 2002. Karyotypic diversity in a *Bryconamericus* aff. *exodon* population (Characidae, Tetragonopterinae). Cytologia. 67(4):397-402.
- Pazza R, Kavalco SAF, Penteado PR, Kavalco KF, DeAlmeida-Toledo LF. 2008. The species complex *Astyanax fasciatus* Cuvier (Teleostei, Characiformes) – a multidisciplinary approach. Journal of Fish Biology. 72(8):2002-2010.
- Pazian MF, Pereira LHG, Shimabukuru-Dias CK, Oliveira C, Foresti F. 2012. Cytogenetic and molecular markers reveal the complexity of the genus *Piabina* Reinhardt, 1867 (Characiformes: Characidae). Neotropical Ichthyology. 10:329-340.
- Peres WAM, Bertollo LAC, Buckup PA, Blanco DR, Kantek DLZ, Moreira-Filho O. 2012. Invasion, dispersion and hybridization of fish associated to river transposition: Karyotypic evidence in *Astyanax "bimaculatus group"* (Characiformes: Characidae). Reviews in Fish Biology and Fisheries. 22:519-526.
- Peres WAM, Buckup PA, Kantek DLZ, Bertollo LAC, Moreira-Filho O. 2009. Chromosomal evidence of downstream dispersal of *Astyanax fasciatus* (Characiformes, Characidae) associated with river shed interconnection. Genetica. 137:305-311.
- Pinkel D, Straume T, Gray JW. 1986. Cytogenetic analysis using quantitative, high-sensitivity, fluorescence hybridization. Proc Natl Acad Sci. 83:2934-2938.
- Piscor D, Paiz LM, Baumgärtner L, Cerqueira FJ, Fernandes CA, Lui RL, Margarido VP. 2020. Chromosomal mapping of repetitive sequences in *Hypessobrycon eques* (Characiformes, Characidae): a special case of the spreading of 5S rDNA clusters in a genome. Genetica. 148(1):25-32.
- Piscor D, Centofante L, Parise-Maltempi PP. 2017. Distinct classical and molecular cytogenetics of *Astyanax marionae* and *A. fasciatus* (Characiformes: Characidae): a comparative study of the organization of heterochromatin and repetitive genes. Journal of genetics. 96:665-671.
- Piscor D, Parise-Maltempi. 2015. First description of B chromosomes in the *Hypessobrycon* (Characiformes, Characidae) genus: a hypothesis for the extra element of *Hypessobrycon eques* Steindachner, 1882. Comparative Cytogenetics. 9(3):325.
- Piscor D, Ribacinko-Piscor DB, Fernandes CA, Parise-Maltempi PP. 2013. Cytogenetic analysis in three *Bryconamericus* species (Characiformes, Characidae): first description of the 5S rDNA-bearing chromosome pairs in the genus. Molecular Cytogenetics. 6:1-8.
- Portela BA, Galetti-Jr. PM, Bertollo LAC. 1988. Considerations on the chromosome evolution of Tetragonopterinae (Pisces, Characidae). Braz. J. Genet. 11(2):307-316.
- Portela-Castro ALB, Júlio-Jr, HF. 2002. Karyotype relationships among species of the subfamily Tetragonopterinae (Pisces, Characidae): Cytotaxonomic and evolution aspects. Cytologia. 67:329-336.
- Prestes AB, Nardelli A, Paiz LM, Gavazzoni M, Margarido VP. 2019. Cytogenetic markers as tools in delimiting species of the highly diverse neotropical fish *Bryconamericus* (Characiformes: Characidae). Neotropical Ichthyology. 17(3):e190057.
- Reis RE, Kullander SO, Ferraris-Jr CJ. 2003. Check list of the freshwater fishes of South and Central America. Porto Alegre: Edipucrs.
- Rubert M, Margarido VP. 2007. Cytogenetic studies in three species of the genus *Oligosarcus*. Brazilian Archives of Biology and Technology. 50:127-135.
- Salvador LB, Moreira-Filho O. 1992. B chromosomes in *Astyanax scabripinnis* (Pisces, Characidae). Heredity. 69(1):50-56.
- Santos ARD, Usso MC, Gouveia JG, Araya-Jaime C, Frantine-Silva W, Giuliano-Caetano L, Dias AL. 2017. Chromosomal mapping of repetitive DNA sequences in the genus *Bryconamericus* (Characidae) and DNA barcoding to differentiate populations. Zebrafish. 14(3):261-271.
- Sánchez KI, Takagui FH, Fenocchio AS. 2021. Cytogenetic analyses in three species of *Moenkhausia* Eigenmann, 1903 (Characiformes, Characidae) from

- Upper Paraná River (Misiones, Argentina). *Caryologia*. 74(4):3-10.
- Sánchez-Romero O, Abad CQ, Cordero PQ, de Sene VF, Nirchio M, Oliveira C. 2015. First description of the karyotype and localization of major and minor ribosomal genes in *Rhoadsia altipinna* Fowler, 1911 (Characiformes, Characidae) from Ecuador. *Comparative cytogenetics*. 9(2):271.
- Scheel JJ. 1973. Fish chromosomes and their evolution. Internal Report of Danmarks Akvarvum, 22.
- Schmid M. 1980. Chromosome banding in amphibia. IV. Differentiation of GC-and AT-rich chromosome regions in Anura. *Chromosoma*. 77(1):83-103.
- Schmid M, Ziegler CG, Steinlein C, Nanda I, Schartl M. 2006. Cytogenetics of the bleak (*Alburnus alburnus*), with special emphasis on the B chromosomes. *Chromosome Research*, 14, 231-242.
- Silva GS, Melo BF, Oliveira C, Benine RC. 2016. Revision of the South American genus *Tetragonopterus* Cuvier, 1816 (Teleostei: Characidae) with description of four new species. *Zootaxa*. 4200(1):1-46.
- Silva DM, Pansonato-Alves JC, Utsunomia R, Daniel SN, Hashimoto DT, Oliveira C, Foresti F. 2013. Chromosomal organization of repetitive DNA sequences in *Astyanax bockmanni* (Teleostei, Characiformes): dispersive location, association and co-localization in the genome. *Genetica*. 141:329-336.
- Silva PC, Santos U, Travenzoli NM, Zanuncio JC, Cioffi MDB, Dergam JA. 2012. The unique karyotype of *Henochilus wheatlandii*, a critically endangered fish living in a fast-developing region in Minas Gerais State, Brazil. *Plos One*. 7(7): e42278.
- Soares LB, Paim FG, Ramos LP, Foresti F, Oliveira C. 2021. Molecular cytogenetic analysis and the establishment of a cell culture in the fish species *Hollandichthys multifasciatus* (Eigenmann & Norris, 1900) (Characiformes, Characidae). *Genetics and Molecular Biology*. 44(2):e20200260.
- Soto MÁ, Castro JP, Walker LI, Malabarba LR, Santos MH, de Almeida MC, Artoni RF. 2018. Evolution of trans-Andean endemic fishes of the genus *Cheirodon* (Teleostei: Characidae) are associated with chromosomal rearrangements. *Revista chilena de historia natural*. 91(1):1-8.
- Sousa CRP, Dos Santos JLA, Silva-Oliveira GC, Furo IDO, de Oliveira EHC, Vallinoto M. 2023. Characterization of a new cytotype and occurrence of a B microchromosome in two spot *Astyanax*, *Astyanax bimaculatus* Linnaeus, 1758 (Characiformes: Characidae). *Journal of Fish Biology*. 102(2):520-524.
- Sumner ATA. 1972. simple technique for demonstrating centromeric heterochromatin. *Exp. Cell Res.* 75:304-306.
- Tenório RCCDO, Vitorino CDA, Souza IL, Oliveira C, Venere PC. 2013. Comparative cytogenetics in *Astyanax* (Characiformes: Characidae) with focus on the cytotaxonomy of the group. *Neotropical Ichthyology*. 11:553-564.
- Terán GE, Benítez MF, Mirande JM. 2020. Opening the Trojan horse: phylogeny of *Astyanax*, two new genera and resurrection of *Psalidodon* (Teleostei: Characidae). *Zoological Journal of the Linnean Society*. 190(4):1217-1234.
- Tonello S, Blanco DR, Cerqueira FJ, Lira NL, Traldi JB, Pavanelli CS, Lui RL. 2022. High rDNA polymorphisms in *Astyanax lacustris* (Characiformes: Characidae): new insights about the cryptic diversity in *A. bimaculatus* species complex with emphasis on the Paraná River basin. *Neotropical Ichthyology*. 20(2):e210147.
- Torres RA, Motta TS, Nardino D, Adam ML, Ribeiro J. 2008. Chromosomes, RAPDs and evolutionary trends of the Neotropical fish *Mimagoniates microlepis* (Teleostei: Characidae: Glandulocaudinae) from coastal and continental regions of the Atlantic Forest, Southern Brazil. *Acta Zoologica*. 89(3):253-259.
- Torres-Mariano AR, Morelli S. 2008. B chromosomes in a population of *Astyanax eigenmanniorum* (Characiformes, Characidae) from the Araguari River Basin (Uberlândia, MG, Brazil). *Genetics and Molecular Biology*. 31:246-249.
- Troy WP, Pacheco ÉB, Oliveira C, Miyazawa CS. 2010. Chromosomal number characterization of *Odontostilbe* and *Serrapinnus* species (Characidae: Cheirodoninae) from Paraguay River basin, Brazil. *Publicatio UEPG-Ciências Biológicas e da Saúde*. 16(1):57-61.
- Ussó MC, Mortati AF, Morales-Blanco AG, Giuliano-Caetano L, Dias AL. 2018. Molecular cytogenetics in different populations of *Oligosarcus paranensis* (Characidae): comparative analysis of the genus with 5S and 18S rDNA probes. *Caryologia*. 71(2):103-108.
- Voltolin DTHTA, de Arruda ADNV, Foresti F, Bortolozzi J, Porto-Foresti F. 2012. Cytogenetic analysis of B chromosomes in one population of the fish *Moenkhausia sanctaefilomenae* (Steindachner, 1907) (Teleostei, Characiformes). *Comparative Cytogenetics*. 6(2):141-151.
- Wasko AP, Galetti Jr PM. 1998. Karyotype diversity in the neotropical fish *Bryconamericus* (Characidae, Tetragonopterinae). *Cytobios*. 94:185-193.
- Wasko AP, Vénere PC, Galetti Jr PM. 1996. Chromosome divergence between two sympatric characid fishes of the genus *Bryconamericus*. *Braz J. Genet.* 19:225-230.
- Yano CF, Semer A, Kretschmer R, Bertollo LAC, Ezaz T, Hatanaka T, de Bello Cioffi M. 2021. Against the

mainstream: exceptional evolutionary stability of ZW sex chromosomes across the fish families Triportheidae and Gasteropelecidae (Teleostei: Characiformes). Chromosome Research. 29(3-4):391-416.

Yano CF, Bertollo LAC, Liehr T, Troy WP, Cioffi MDB. 2016. W chromosome dynamics in *Triportheus* spe-

cies (Characiformes, Triportheidae): an ongoing process narrated by repetitive sequences. Journal of Heredity. 107(4):342-348.

Table 1. Cytogenetic data of small characins and closely related groups, according to Mirande (2018) and Terán *et al.* (2020) (the asterisks indicate the taxa whose scientific names were updated).

Species	2n	Karyotype	Locality	Distinctive large m pair	Reference
Family Characidae					
Subfamily Aphyocharacinae					
<i>Prionobrama filigera</i>	52	12m/sm+ 40st/a	Not Informed	Present	Arefjev (1990)
Subfamily Characinae					
<i>Phenacogaster cf. pectinatus</i>	46	12m+2st+32a	São Francisco stream - AC	Present	Carvalho <i>et al.</i> (2002)
Subfamily Cheirodontinae					
<i>Cheirodon australis</i>	52	8m+6sm+4st+33a	La Poza Lake - Chile	Present	Soto <i>et al.</i> (2018)
<i>Cheirodon galusdae</i>	52	6m+6sm+4st+34a	Andalién River - Chile	Present	Soto <i>et al.</i> (2018)
<i>Cheirodon interruptus</i>	52	6m+6sm+4st+34a	Marga-Marga River - Chile	Present	Soto <i>et al.</i> (2018)
<i>Cheirodon kiliani</i>	52	8m+6sm+4st+33a	Calle-Calle River - Chile	Present	Soto <i>et al.</i> (2018)
<i>Cheirodon pisciculus</i>	52	6m+6sm+4st+34a	Angostura River - Chile	Present	Soto <i>et al.</i> (2018)
<i>Odontostilbe pequira</i>	52	14m+20sm+14st+4a	Onça stream - MS	Absent	Nishiyama <i>et al.</i> (2015)
<i>Odontostilbe pequira</i>	52	24m+12sm+12st+4a	Cuiabá River - MT	Absent	Troy <i>et al.</i> (2010)
<i>Serrapinnus calliurus</i>	52	36m+12sm+6st	Bento Gomes River - MT	Absent	Troy <i>et al.</i> (2010)
<i>Serrapinnus heterodon</i>	52	16m+20sm+14st+2a	São Francisco River - MG	Absent	Peres <i>et al.</i> (2007)
<i>Serrapinnus kriegi</i>	52	24m+18sm+10st	Cuiabá River - MT	Absent	Troy <i>et al.</i> (2010)
<i>Serrapinnus microdon</i>	52	30m+12sm+8st+4a	Bento Gomes River - MT	Absent	Troy <i>et al.</i> (2010)
<i>Serrapinnus piaba</i>	52	16m+20sm +14st+2a	São Francisco River- MG	Absent	Peres <i>et al.</i> (2007)
Subfamily Stethaprioninae					
<i>Astyanax abramis</i>	50	4m+30sm+8st+8a	Iguaçu River - PR	Present	Gavazzoni <i>et al.</i> (2018)
<i>Astyanax altiparanae</i>	50	6m+28sm+4st+12a	Pântano Stream and Jordão River - PR, Feijão Stream - MG	Present	Ferreira-Neto <i>et al.</i> (2009)
<i>Astyanax altiparanae</i>	50	6m+28sm+8st+8a	Tibagi River - PR	Present	Domingues <i>et al.</i> (2007)
<i>Astyanax altiparanae</i>	50	6m+30sm+8st+6s	Iguaçu River - PR	Present	Domingues <i>et al.</i> (2007)
<i>Astyanax altiparanae</i>	50	16m+24sm+4st+6a	Queixada River - PR	Present	Da Silva <i>et al.</i> (2016)
<i>Astyanax altiparanae</i>	50	16m+20sm+4st+10a	Esperança stream - PR	Present	Da Silva <i>et al.</i> (2016)
<i>Astyanax altiparanae</i>	50	16m+20sm+4st+10a	Jacutinga River - PR	Present	Da Silva <i>et al.</i> (2016)
<i>Astyanax altiparanae</i>	50	6m+28sm+4st+12a	Paraná River - PR	Present	Gavazzoni <i>et al.</i> (2018)
<i>Astyanax asuncionensis</i>	50	18m+22sm+6st+4a	Miranda River - MS	Present	Da Silva <i>et al.</i> (2016)
<i>Astyanax asuncionensis</i>	50	8m+24sm+6st	Iguaçu River - PR	Present	Gavazzoni <i>et al.</i> (2018)
<i>Astyanax aff. bimaculatus</i>	50	4m+14sm+24st+8a	Dois de Agosto stream - PA	Present	Sousa <i>et al.</i> (2023)
<i>Astyanax bimaculatus</i>	50	8m+34sm+2st+6a	São Francisco River - PR	Present	Peres <i>et al.</i> (2012)
<i>Astyanax bimaculatus</i>	50	8m+32sm+2st+8a	Grande River - PR	Present	Peres <i>et al.</i> (2012)
<i>Astyanax bimaculatus</i>	50	8m+33sm+2st+7a	Piumhi River - PR	Present	Peres <i>et al.</i> (2012)
<i>Astyanax bimaculatus</i>	50	8m+31sm+2st+9a	Piumhi River - PR	Present	Peres <i>et al.</i> (2012)
<i>Astyanax bimaculatus</i>	50	8m+30sm+2st+10a	Piumhi River - PR	Present	Peres <i>et al.</i> (2012)
<i>Astyanax bimaculatus</i>	50	10m+18sm+12 st+10a	Aguapeí River - SP	Present	Alberdi and Fenocchio (1997)
<i>Astyanax bimaculatus</i>	50	6m+28sm+8st+8a	Caeté River - PA	Present	Sousa <i>et al.</i> (2023)
<i>Astyanax lacustris</i>	50	10m+24sm+6st+10a	Itaipu Lake, Paraná River basin - PR	Present	Tonello <i>et al.</i> (2022)

Species	2n	Karyotype	Locality	Distinctive large m pair	Reference
<i>Astyanax lacustris</i>	50	6m+12sm+14st+18a	Pirassununga River - SP	Present	Goes <i>et al.</i> (2020)
<i>Astyanax jacuhiensis</i>	50	10m+26sm+6st+8a	Tramandaí River basin - RS	Present	Da Silva <i>et al.</i> (2012)
<i>Astyanax jacuhiensis</i>	50	8m+30sm+4st+8a	Guaíba Lake - RS	Present	Pacheco <i>et al.</i> (2010)
<i>Astyanax jacuhiensis</i>	50	8m+28sm+6st+8a	Ijuá River - PR	Present	Gavazzoni <i>et al.</i> (2018)
<i>Astyanax scabripinnis</i>	50	8m+20s+8st+14a	Macacos River - PR	Present	Kavalco and Moreira-Filho (2003)
<i>Astyanax scabripinnis</i>	50	6m+22sm+10st+12a	Córrego das Pedras stream - SP	Present	Mestriner <i>et al.</i> (2000)
<i>Astyanax scabripinnis</i>	50	8m+22sm+12st+6a	Mogi-Guaçu River basin - SP	Present	Pazza <i>et al.</i> (2008)
<i>Astyanax scabripinnis</i>	50	12m+20sm+10st+4a	Paranapanema River basin - SP	Present	Pazza <i>et al.</i> (2008)
<i>Astyanax scabripinnis</i>	50	6m+22sm+10st+12a	Córrego das Pedras stream - SP	Present	Salvador and Moreira-Filho (1992)
<i>Astyanax scabripinnis</i>	50	10m+20sm+8st+12a	São Francisco River - PR	Present	Klassmann and Martins-Santos (2017)
<i>Astyanax scabripinnis</i>	48	11m+18sm+9st+10a	São Francisco River - PR	Present	Klassmann and Martins-Santos (2017)
<i>Astyanax scabripinnis</i>	48	10m+20sm+8st+10a	Ivaí River - PR	Present	Alves and Martins-Santos (2002)
<i>Astyanax</i> sp.	50	4m+22sm+8st+16a	Piraquara, Upper Paraná River basin - PR	Present	Kantek <i>et al.</i> (2008)
<i>Astyanax</i> sp.	50	4m+22sm+8st+16a	Bicudo River, Upper Paraná River basin - PR	Present	Kantek <i>et al.</i> (2008)
<i>Astyanax</i> sp.	50	4m+24sm+6st+16a	Bicudo River, Upper Paraná River basin - PR	Present	Kantek <i>et al.</i> (2008)
<i>Astyanax</i> sp.	52	22m+26sm+4a	Upper Paraná River basin - PR	Absent	Tenório <i>et al.</i> (2013)
<i>Brachychalcinus retrospina</i>	50	6m+24sm+6st+4a	Angelim River - MT	Present	Krinski and Miyazawa (2012)
<i>Ctenobrycon hauxwellianus</i>	50	10m+6sm+34st	São Francisco stream - AC	Present	Carvalho <i>et al.</i> (2002)
<i>Deuterodon (Astyanax) giton*</i>	50	6m+8sm+8st+28a	Paraitinga River - SP	Present	Kavalco and Moreira-Filho (2003)
<i>Deuterodon (Astyanax) intermedius*</i>	50	6m+18sm+12st+10a	Paraitinga River - SP	Present	Kavalco and Moreira-Filho (2003)
<i>Deuterodon (Astyanax) janeiroensis*</i>	50	6m+14sm+14st+16a	Betari River - SP	Present	Carvalho <i>et al.</i> (2002)
<i>Deuterodon pedri</i>	50	12m+12sm+20st+6a	Santo Antônio - River	Present	Coutinho-Sanches and Dergam (2015)
<i>Deuterodon pedri</i>	50	14m/sm+36st/a	Pedri River - SP	Present	Portela <i>et al.</i> (1988)
<i>Deuterodon stigmaturus</i>	50	8m+6sm+2st+34a	Maquiné River - RS	Present	Mendes <i>et al.</i> (2011)
<i>Gymnocorhimbis ternetzi</i>	50	14m+12sm+6st	Paraná River - PR	Present	Alberdi and Fenocchio (1997)
<i>Hasemania crenuchoides</i>	50	6m+26sm+16st+2a	Alto-Tocantins River	Present	Duarte <i>et al.</i> (2018)
<i>Hasemania marginata</i>	50	12m+18sm+10st+10a.	Not Informed	Present	Arefjev (1990)
<i>Hasemania nana</i>	50	8m+42sm	São Francisco River basin - MG	Present	Moreira <i>et al.</i> (2007)
<i>Hemigrammus hyanuary</i>	52	22m/sm+30st/a.	Not Informed	Present	Arefjev (1990)
<i>Hemigrammus marginatus</i>	50	10m+34sm+6a	Upper Paraná River basin - PR	Present	Portela-Castro and Júlio-Jr (2002)
<i>Hollandichthys multifasciatus</i>	50	8m+10sm+32st	Iguapé River - SP	Present	Soares <i>et al.</i> (2021)
<i>Hollandichthys multifasciatus</i>	50	10m+12sm+28st	Grande River - SP	Present	Carvalho <i>et al.</i> (2002)
<i>Hypessobrycon anisitsi</i>	50	6m+16sm+12st+16a	Upper Paraná River - PR	Present	Centofante <i>et al.</i> (2003)
<i>Hypessobrycon anisitsi</i>	50	10m+2sm+20st+18a	Pirassununga River -SP	Present	Goes <i>et al.</i> (2020)
<i>Hypessobrycon anisitsi</i>	50	18m+10sm+6st+16a	Tibagí River - PR	Present	Mendes <i>et al.</i> (2011)
<i>Hypessobrycon eques</i>	52	10m+20sm+8st+14a	Piracicaba River - SP	Absent	Piscor <i>et al.</i> (2020)
<i>Hypessobrycon eques</i>	52	14m+16sm+4st+18a	Capivara River - SP	Absent	Martinez <i>et al.</i> (2012)
<i>Hypessobrycon eques</i>	52	12m+26sm+8st+14a	Ribeirão Claro River - SP	Absent	Piscor and Parise-Maltempi (2015)
<i>Hypessobrycon flammeus</i>	52	18m/sm+32st+2a.	Not informed	Present	Arefjev (1990)
<i>Hypessobrycon herbertaxelrodi</i>	52	10m/sm+42st/a.	não informado	Absent	Arefjev (1990)
<i>Hypessobrycon luetkenii</i>	50	6m + 8sm + 36a	Maquiné River - RS	Present	Mendes <i>et al.</i> (2011)
<i>Hypessobrycon reticulatus</i>	50	14m+20sm+16st	Jequiá River - SP	Present	Carvalho <i>et al.</i> (2002)

Species	2n	Karyotype	Locality	Distinctive large m pair	Reference
<i>Hyphessobrycon scholzei</i>	50	8m+20sm+14a	Not Informed	Present	Arefjev (1990)
<i>Hyphessobrycon vinaceus</i>	50	8m+12sm+30a	Pardo River - BA	Present	Nishiyama <i>et al.</i> (2015)
<i>Moenkhausia cosmops</i>	50	14m+30sm+6st	Verde River - MT	Present	Nascimento <i>et al.</i> (2020)
<i>Moenkhausia costae</i>	50	50m/sm	São Francisco River - MG	Present	Portela <i>et al.</i> (1988)
<i>Moenkhausia dichroura</i>	50	22m+22sm+6st	Upper Paraná River - Argentina	Present	Sánchez <i>et al.</i> (2021)
<i>Moenkhausia forestii</i>	50	10m+32sm+8st	Sangue River - MT	Present	Nascimento <i>et al.</i> (2020)
<i>Moenkhausia intermedia</i>	50	16m+28sm+6st	Upper Paraná River - Argentina	Present	Sánchez <i>et al.</i> (2021)
<i>Moenkhausia intermedia</i>	50	50m/sm	Lagoa do Mato - SP	Present	Portela <i>et al.</i> (1988)
<i>Moenkhausia intermedia</i>	50	16m+34sm	Paraná River - PR	Present	Portela-Castro and Júlio-Jr (2002)
<i>Moenkhausia nigromarginata</i>	50	14m+32sm+4a	Verde River - MT	Present	Nascimento <i>et al.</i> (2020)
<i>Moenkhausia oligolepis</i>	50	12m+32sm+6st	Xapuri River- AC	Present	Nascimento <i>et al.</i> (2020)
<i>Moenkhausia sanctaefilomenae</i>	50	6m+23sm+12st	Batalha River, Tietê River basin - SP	Present	Voltolin <i>et al.</i> (2012)
<i>Moenkhausia sanctaefilomenae</i>	50	48m/sm+2a	Capivara and Tietê River - SP	Present	Forestii <i>et al.</i> (1989)
<i>Moenkhausia sanctaefilomenae</i>	50	12m+36sm+2a	Paraná River - PR	Present	Portela-Castro and Júlio-Jr (2002)
<i>Moenkhausia sanctaefilomenae</i>	50	12m+32sm+6st	Upper Paraná River - Argentina	Present	Sánchez <i>et al.</i> (2021)
<i>Moenkhausia sanctaefilomenae</i>	50	48m/sm+2st/a	Aguapeí River - Argentina	Present	Alberdi and Fenocchio (1997)
<i>Nematabrycon palmeri</i>	50	8m/sm+10st+32a	Not Informed	Present	Arefjev (1990)
<i>Nematocharax venustus</i>	50	8m+26sm+14st+2a	Contas River - BA/Jequitinhonha - MG	Present	Barreto <i>et al.</i> (2016)
<i>Oligosarcus acutirostris</i>	50	4m+14sm+18st	Paraibuna River - ES	Present	Cunha <i>et al.</i> (2021)
<i>Oligosarcus hepsetus</i>	50	6m+10sm+16+18a	Paraitinga and Paraíba do Sul River basin -SP	Present	Kavalco <i>et al.</i> (2005)
<i>Oligosarcus hepsetus</i>	50	2m+16sm+16st+16a	Paraíba do Sul River - SP	Present	Hattori <i>et al.</i> (2007)
<i>Oligosarcus jenynsii</i>	50	2m+24sm+10st+14a	Uruguay River - SC	Present	Hattori <i>et al.</i> (2007)
<i>Oligosarcus longirostris</i>	50	4m+10sm+16st+20a	Iguaçu River - PR	Present	Rupert and Margarido (2007)
<i>Oligosarcus paranensis</i>	50	8m+18sm+10st+14a	Tibagi River basin - PR	Present	Ussó <i>et al.</i> (2018)
<i>Oligosarcus paranensis</i>	50	4m+10sm+16st+20a	Piquiri River basin - PR	Present	Rupert and Margarido (2007)
<i>Oligosarcus pintoi</i>	50	4m+10sm+16st+20a	Piquiri River basin - PR	Present	Rupert and Margarido (2007)
<i>Oligosarcus pintoi</i>	50	4m+12sm +14st +20a	Ivaí River - PR	Present	Mari-Ribeiro <i>et al.</i> (2022)
<i>Oligosarcus pintoi</i>	50	2m+20sm+12st+16a	Mogi-Guaçu River - SP	Present	Hattori <i>et al.</i> (2007)
<i>Oligosarcus</i> sp.	50	6m+14sm+18st+12a	Velhas River basin - Ouro Preto - MG	Present	De-Barros <i>et al.</i> (2015)
<i>Oligosarcus</i> sp.	50	4m+14sm+20st+12a	Doce River basin - MG	Present	De-Barros <i>et al.</i> (2015)
<i>Orthopinus franciscensis</i>	50	22m+20sm+2st+6a	São Francisco River - MG	Present	Moreira <i>et al.</i> (2007)
<i>Poptella paraguayensis</i>	50	10m+26sm+8st+6a	Miranda River - MT	Present	Freitas and Galetti (1998)
<i>Psalidodon (Astyanax) bockmanni*</i>	50	10m+12sm+12st+16a	Paranapanema River basin, São Miguel Arcanjo and Pilar do Sul - SP	Present	Kavalco <i>et al.</i> (2008)
<i>Psalidodon (Astyanax) bockmanni*</i>	50	8m+14sm+12st+16a	Capivara River, Tietê River basin - SP	Present	Silva <i>et al.</i> (2013)
<i>Psalidodon (Astyanax) bockmanni*</i>	50	8m+14sm+14st+14a	Água Madalena stream, Paranapanema River basin - SP	Present	Silva <i>et al.</i> (2013)
<i>Psalidodon (Astyanax) bockmanni*</i>	50	6m+20sm+8st+16a	Iguatemi River basin - MS	Present	Fernandes <i>et al.</i> (2010)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	50	16m+12sm+6st+16a	Tributary of Cabeça River - SP	Present	Piscor <i>et al.</i> (2017)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	48	10m+20sm+8st+10a	Tributary of Ribeirão Claro River - SP	Present	Piscor <i>et al.</i> (2017)
<i>Psalidodon (Astyanax) eigenmanniorum*</i>	48	14m+24sm+4st+10a	Araguari River Basin - MG	Present	Torres-Mariano and Morelli (2008)
<i>Psalidodon (Astyanax) eigenmanniorum*</i>	48	10m+16sm+10st+12a	Laguna dos patos - RS	Present	Mendes <i>et al.</i> (2011)

Species	2n	Karyotype	Locality	Distinctive large m pair	Reference
<i>Psalidodon (Astyanax) fasciatus*</i>	48	8m+20sm+16st+4a	São Francisco River - MG	Present	Peres <i>et al.</i> (2009)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	50	8m+26sm+6st+10a	Afluente do rio Corumbataí - SP	Present	Piscor <i>et al.</i> (2017)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	48	10m+12sm+12st+14a	Pirassununga - SP	Present	Goes <i>et al.</i> (2020)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	48	8m+20sm+12st+8a	Preto do Costa River - BA	Present	Medrado <i>et al.</i> (2015)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	48	8m+18sm+14st+8a	Mutum River- BA	Present	Medrado <i>et al.</i> (2015)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	48	8m+24sm+10st+6a	Oricó River - BA	Present	Medrado <i>et al.</i> (2015)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	48	8m+28sm+8st+4a	Criciúma River - BA	Present	Medrado <i>et al.</i> (2015)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	48	8m+18sm+16st+6a	Gongogi River - BA	Present	Medrado <i>et al.</i> (2015)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	48	8m+16sm+16st+8a	Mineiro River - BA	Present	Medrado <i>et al.</i> (2015)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	48	8m+16sm+18st+6a	Itapicuru River - BA	Present	Medrado <i>et al.</i> (2015)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	48	8m+24sm+10st+6a	Braço River - BA	Present	Medrado <i>et al.</i> (2015)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	48	8m+22sm+10st+8a	Cachoeira River - BA	Present	Medrado <i>et al.</i> (2015)
<i>Psalidodon (Astyanax) aff. fasciatus*</i>	48	8m+20sm+16st+4a	Contas River - BA	Present	Medrado <i>et al.</i> (2015)
<i>Psalidodon (Astyanax) marionae*</i>	48	4m+24sm+10st+6a	Rio claro stream, Paraguai River basin - MS	Present	Piscor <i>et al.</i> (2017)
<i>Psalidodon (Astyanax) parabybae*</i>	48	8m+18sm+12st+10a	Paraitinga River - SP	Present	Kavalco and Moreira-Filho (2003)
<i>Psalidodon (Astyanax scabripinnis paranae) paranae*</i>	50	4m+34sm+4st+6a	Araquá River - SP	Present	Maistro <i>et al.</i> (1992)
<i>Psalidodon (Astyanax) schubarti*</i>	36	10m+10sm+10st+6a	Pirassununga -SP	Present	Goes <i>et al.</i> (2020)
<i>Psalidodon (Astyanax) schubarti*</i>	36	14m+14sm/6st+2a	Paraná River - PR	Present	Alberdi and Fenocchio (1997)
<i>Rhoadsia altipinna</i>	50	10m+26+14a	Das Bocas River - Ecuador	Present	Sanchez-Romero <i>et al.</i> (2015)
Subfamily Stevardiinae					
<i>Bryconamericus aff. exodon</i>	52	16m+12sm+6st+18a	Tibagí River - PR	Absent	Paintner-Marques <i>et al.</i> (2002)
<i>Bryconamericus aff. exodon</i>	52	10m+24sm+6st+12a	Tibagí River - PR	Absent	Paintner-Marques <i>et al.</i> (2002)
<i>Bryconamericus aff. iheringii</i>	52	12m+10sm+16st+14a	Três Bocas Stream - PR	Absent	Da Silva <i>et al.</i> (2014)
<i>Bryconamericus aff. iheringii</i>	52	18m+14sm+10st+10a	Três Bocas Stream - PR	Absent	Da Silva <i>et al.</i> (2014)
<i>Bryconamericus aff. iheringii</i>	52	20m+18sm+4st+10a	Três Bocas Stream - PR	Absent	Da Silva <i>et al.</i> (2014)
<i>Bryconamericus aff. iheringii</i>	52	20m+14sm+12st+6a	Três Bocas Stream - PR	Absent	Da Silva <i>et al.</i> (2014)
<i>Bryconamericus aff. iheringii</i>	52	22m+18sm+8st+4a	Três Bocas Stream - PR	Absent	Da Silva <i>et al.</i> (2014)
<i>Bryconamericus aff. iheringii</i>	52	18m+24sm+6st+4a	Três Bocas Stream - PR	Absent	Da Silva <i>et al.</i> (2014)
<i>Bryconamericus aff. iheringii</i>	52	12m+18sm+8st+ 14a	Maringá stream, Paraná River basin - PR	Absent	Capistano <i>et al.</i> (2008)
<i>Bryconamericus aff. iheringii</i>	52	8m+28sm+6st+ 10a	Keller River, Paraná River basin - PR	Absent	Capistano <i>et al.</i> (2008)
<i>Bryconamericus aff. iheringii</i>	52	8m+20sm+8st+16a	Tatupeba stream, Paraná River basin - PR	Absent	Capistano <i>et al.</i> (2008)
<i>Bryconamericus aff. iheringii</i>	52	10m+16sm+14st+12a	Upper Uruguai River basin	Present	Prestes <i>et al.</i> (2009)

Species	2n	Karyotype	Locality	Distinctive large m pair	Reference
<i>Bryconamericus</i> aff. <i>iheringii</i>	52	12m+18sm+8st+14a	Ocoí River - PR	Absent	Nishiyama <i>et al.</i> (2015)
<i>Bryconamericus</i> aff. <i>iheringii</i>	52	10m+14sm+18st+10a	Corumbataí River - SP	Absent	Piscor <i>et al.</i> (2013)
<i>Bryconamericus coeruleus</i>	52	14m+20sm+8st+10a	Upper Paraná River basin - PR	Present	Prestes <i>et al.</i> (2009)
<i>Bryconamericus ecai</i>	52	10m+16sm+8st +18a	Forquetinha River - RS	Absent	Santos <i>et al.</i> (2017)
<i>Bryconamericus ecai</i>	52	8m+16sm+14st+14a	Forquetinha River - RS	Absent	Santos <i>et al.</i> (2017)
<i>Bryconamericus ecai</i>	52	10m+16sm+8st+18a	Forquetinha River - RS	Absent	Santos <i>et al.</i> (2017)
<i>Bryconamericus ecai</i>	52	10m+10sm+8st+24a	Forquetinha River - RS	Absent	Dos Santos <i>et al.</i> (2012)
<i>Bryconamericus ecai</i>	52	10m+18sm+8st+16a	Forquetinha River - RS	Absent	Dos Santos <i>et al.</i> (2012)
<i>Bryconamericus ecai</i>	52	14m+14sm+6st+18a	Forquetinha River - RS	Absent	Dos Santos <i>et al.</i> (2012)
<i>Bryconamericus ecai</i>	52	10m+24sm+14st+4a	Forquetinha River - RS	Absent	Dos Santos <i>et al.</i> (2012)
<i>Bryconamericus ecai</i>	52	10m+16sm+14st+12a	Upper Uruguay River basin - PR	Absent	Prestes <i>et al.</i> (2009)
<i>Bryconamericus eigenmanni</i>	52	6m+16sm+16st+14a	Upper Uruguay River basin - PR	Present	Prestes <i>et al.</i> (2009)
<i>Bryconamericus</i> sp.	52	16m+14sm+10st+12a	Vermelho stream, Ivaí River basin - PR	Absent	Santos <i>et al.</i> (2017)
<i>Bryconamericus</i> sp.	52	2m+12sm+20st+20a	Cambuta River, Ivaí River basin - PR	Absent	Santos <i>et al.</i> (2017)
<i>Bryconamericus</i> sp. A	52	6m+30sm+6st+10a	Piracicaba river - SP	Absent	Wasko and Galetti-Jr (1998)
<i>Bryconamericus</i> sp. B	52	6m+10sm+20st+16a	Piracicaba river - SP	Absent	Wasko and Galetti-Jr (1998)
<i>Bryconamericus</i> sp. C	52	6m+18sm+14st+14a	Tibagi River - PR	Absent	Wasko and Galetti-Jr (1998)
<i>Bryconamericus</i> sp. D	52	8m+14sm+16st+14a	Garças River - MT	Absent	Wasko and Galetti-Jr (1998)
<i>Bryconamericus</i> sp. E	54	10m+16sm+22st+6a	Garças River - MT	Absent	Wasko and Galetti-Jr (1998)
<i>Bryconamericus</i> sp. A	52	6m+30sm+ 6st+10a	Piracicaba river - SP	Absent	Wasko <i>et al.</i> (1996)
<i>Bryconamericus</i> sp. B	52	10m+6sm+18st+18a	Piracicaba river - SP	Absent	Wasko <i>et al.</i> (1996)
<i>Bryconamericus stramineus</i>	52	26m/sm+26st/a	Mogi Guaçu River - SP	Absent	Portela <i>et al.</i> (1988)
<i>Bryconamericus stramineus</i>	52	6m+10sm+16st+20a	Iguatemi River basin - MS	Absent	Fernandes <i>et al.</i> (2010)
<i>Bryconamericus stramineus</i>	52	6m+10sm+16st+20a	Guaçu stream, Iguatemi River basin - MS	Absent	Piscor <i>et al.</i> (2013)
<i>Bryconamericus turiuba</i>	52	8m+10sm+14st+20a	Passo-Cinco River - SP	Absent	Piscor <i>et al.</i> (2013)
<i>Glandulocauda melanogenys</i>	52	4m+12sm+22st+14a	Paranapiacaba - SP	Absent	Guimarães <i>et al.</i> (1995)
<i>Knodus</i> cf. <i>chapadae</i>	52	14m+14sm+ 14st+10a	Tangará da Serra - MT	Absent	Krinski <i>et al.</i> (2008)
<i>Markiana nigripinnis</i>	52	8m+22sm+22st/a	Miranda River - MT	Absent	Monteiro <i>et al.</i> (2022)
<i>Mimagoniates laterallis</i>	52	6m+20sm+16st+10a	Itanhém - SP	Absent	Guimarães <i>et al.</i> (1995)
<i>Mimagoniates microlepis</i>	52	12m+18sm+14+8a	Iguáçu River basin and Piraquara River - PR	Absent	Torres <i>et al.</i> (2008)
<i>Piabina anhembi</i>	52	8m+10sm+16st+18a	Salesópolis - SP	Absent	Pazian <i>et al.</i> (2012)
<i>Piabina argentea</i>	52	26m/sm+26st/a	Mogi Guaçu - SP	Absent	Pazian <i>et al.</i> (2012)
<i>Piabina argentea</i>	52	8m+14sm+16st+14a	São Francisco - MG	Absent	Pazian <i>et al.</i> (2012)
<i>Piabina argentea</i>	52	4m+22sm+10s+16a	Itatinga - SP	Absent	Pazian <i>et al.</i> (2012)
<i>Piabina argentea</i>	52	8m+18sm+18st+10a	Botucatu - SP	Absent	Pazian <i>et al.</i> (2012)
<i>Piabina argentea</i>	52	4m+24sm+10st+14a	Bauru - SP	Absent	Pazian <i>et al.</i> (2012)
<i>Piabina argentea</i>	52	26m/sm+26st/a	Mogi Guaçu River - SP	Absent	Portela <i>et al.</i> (1988)
<i>Piabina argentea</i>	52	8m+14sm+16st+14a	São Francisco River - MG	Absent	Moreira <i>et al.</i> (2007)
<i>Piabina argentea</i>	52	6m+24sm+12st+10a	Iguatemi River - MS	Absent	Fernandes <i>et al.</i> (2010)
Subfamily Tetragonopterinae					
<i>Tetragonopterus argenteus</i>	52	16m+4sm+4st+28a	Cuiabá River - MT	Absent	Miyazawa (2015)
<i>Tetragonopterus argenteus</i>	52	24m+8sm+4st+16a	Bento Gomes River - MT	Absent	Miyazawa (2015)
<i>Tetragonopterus argenteus</i>	50	14m+4sm+4st+28a	Cuiabá River - MT	Absent	Miyazawa (2015)
<i>Tetragonopterus argenteus</i>	52	16m/sm+2st+34a	Paraná River - PR	Absent	Alberdi and Fenocchio (1997)
<i>Tetragonopterus franciscoensis</i>	52	12m+26sm+14a	Itapicuru River - BA	Absent	Present study
<i>Tetragonopterus franciscoensis</i> (<i>chalceus</i>)*	52	13m/sm+13st/a	São Francisco River - MG	Absent	Portela <i>et al.</i> (1988)

Species	2n	Karyotype	Locality	Distinctive large m pair	Reference
Family Bryconidae					
<i>Brycon amazonicus</i>	50	22m+14sm+14st	Orinoco basin -Venezuela	Present	Mariguela <i>et al.</i> (2010)
<i>Brycon cf. cephalus</i>	50	26m+24sm/st	Amazon basin - AM	Present	Almeida-Toledo <i>et al.</i> (1996)
<i>Brycon cf. reinhardti</i>	50	22m+28sm/st	Paraíba do Sul River - SP	Present	Almeida-Toledo <i>et al.</i> (1996)
<i>Brycon insignis</i>	50	24m+21sm/st	Paraíba do Sul River - SP	Present	Almeida-Toledo <i>et al.</i> (1996)
<i>Henochilus wheatlandii</i>	50	26m+12sm+12st	Santo Antônio River - MG	Present	Silva <i>et al.</i> (2012)
Family Gasteropelecidae					
<i>Carnegiella strigata</i>	50	Not Informed	Manaus - MA	Absent	Yano <i>et al.</i> (2021)
<i>Gasteropelecus levis</i>	54	Not Informed	Manaus - MA	Absent	Yano <i>et al.</i> (2021)
<i>Thoracocharax stellatus</i>	54	Not Informed	Barra do Bugres - MT	Absent	Yano <i>et al.</i> (2021)
Family Triportheidae					
<i>Agoniates halecinus</i>	52	Not Informed	Manaus - AM	Absent	Yano <i>et al.</i> (2021)
<i>Lignobrycon myersi</i>	52	28m+18sm+6a	Almada River - BA	Absent	Dos-Santos <i>et al.</i> (2016)
<i>Triportheus auritus</i>	52	Not Informed	Ponta do Araguaia - MT	Absent	Yano <i>et al.</i> (2021)
<i>Triportheus nematurus</i>	52	13m+23sm+16st	Piracicaba River - SP	Absent	Diniz <i>et al.</i> (2008)
<i>Triportheus pantanensis</i>	52	Not Informed	Paraguay basin	Absent	Yano <i>et al.</i> (2016)
<i>Triportheus aff. rotundatus</i>	52	Not Informed	Paraguay basin	Absent	Yano <i>et al.</i> (2016)