

Evolutionary aspects of the ZZ/ZW sex chromosome system in the Characidae fish, genus *Triporthesus*. A monophyletic state and NOR location on the W chromosome

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Four species/populations of *Triporthesus*, *T. guentheri*, *T. cf. elongatus* and *T. paranense* from different Brazilian hydrographic basins, were studied cytogenetically. All the species showed a similar karyotypic macrostructure, with a diploid chromosome number $2n = 52$ and a ZZ/ZW sex chromosome system. Besides silver- and fluorochrome-staining, the chromosome mapping of 18S rDNA was also investigated using a biotinylated probe. In spite of some variation in the number of the NORs, a major chromosome site was always present on the short arm of an autosomal pair. In addition, a

characteristic rDNA site was also observed on the telomeric region of the W chromosome in the four species/populations. In *Triporthesus* differential reduction in size and heterochromatin accumulation appear to be the main processes associated with the evolution of the sex W chromosome. The location of rRNA genes on this chromosome may correspond to a plesiomorphic condition in the genus and, if so, predates to the sex chromosome system differentiation, with a possible influence in the initial steps of this process. *Heredity* (2002) 89, 15–19. doi:10.1038/sj.hdy.6800081

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Introduction

Triporthesus is a genus of neotropical Characidae fish showing particular features in respect to sex chromosomes. Falcão (1988), described a ZZ/ZW sex chromosome system in four Brazilian species, three of them (*T. elongatus*, *T. flavus* and *T. albus*) belonging to the Amazon basin and the other (*T. signatus*) belonging to the Paraná basin. In all the species, the W chromosome was characterized by a large amount of heterochromatin and a reduced size in relation to the Z chromosome.

After this initial study, additional analyses have also shown such a sex chromosome system in other *Triporthesus* species. This is the case of *T. guentheri* from the São Francisco river, Brazil, which had variation in the size of the W chromosome due to its amount of heterochromatin (Bertollo and Cavallaro, 1992) and *T. paranense* from the Paraná basin, Argentina (Sánchez and Jorge, 1999).

The distribution and differentiation of the ZZ/ZW system in *Triporthesus* were recently reviewed by Artoni *et al* (2001), adding three new reports from different Brazilian hydrographic basins to the previous studies: *T. cf. elongatus* from the Araguaia river, *T. paranense* from Cuiabá river, and *T. paranense* from Paraguai river. Thus, all the species already analyzed (about 50% of the known *Tripor-*

theus species) share the same sex chromosome system, which seems to be a fixed character in the genus. In addition, this ZZ/ZW sex system contrasts with those described for other groups of fish, such as *Leporinus*, *Semaprochilodus* and *Parodon*, in which the W chromosome is enlarged in size due to increased heterochromatin (Galetti Jr and Foresti, 1986; Feldberg *et al*, 1987; Moreira-Filho *et al*, 1993).

In this work the occurrence of rDNA sites on the W sex chromosome of *Triporthesus* was verified, using fluorescence 'in situ' hybridization with a 18S rDNA probe. A possible implication of this rDNA in the evolutionary process of the ZZ/ZW sex chromosome system is also discussed.

Materials and methods

Samples and chromosome preparation

Seventy-four wild individuals of *Triporthesus* fish from different regions of Brazil were analyzed: 20 male and four female *T. guentheri* from the São Francisco river (Três Marias, Minas Gerais State), nine male and two female *T. cf. elongatus* from the Araguaia river (Barra do Garças, Mato Grosso State), 10 male and 24 female *T. paranense* from the Cuiabá river (Cuiabá, Mato Grosso State) and three male and two female *T. paranense* from the Paraguai river (Corumbá, Mato Grosso do Sul State).

Chromosome spreads were obtained by direct preparations from kidney cells (Bertollo *et al*, 1978) and by

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short-term culture of cells (Fenocchio *et al.*, 1991), using stimulation of mitotic activity (Lee and Elder, 1980).

C-banding and Ag-staining

Constitutive heterochromatin (C-banding) was detected according to Sumner (1972), with the following modifications: 2-day-old slides were immersed in 0.2 N HCl for 15 min, and in 5% barium hydroxide at 42°C for 1 min 30 sec. The slides were then quickly washed in 0.2 HCl and incubated in $2 \times$ SSC at 60°C for 60 min, followed by Giemsa staining (5%) for 5 min. After these five steps, slides were washed in deionized water and air-dried. Some preparations were also stained with the fluorochrome DAPI after the conventional C-banding method (C-DAPI), according to Souza and Moreira-Filho (1995).

Nucleolus organizing regions were detected by silver nitrate staining (Ag-NORs), according to Howell and Black (1980). Briefly, the slides were treated with a gelatin-silver nitrate solution, covered with a coverslip, incubated at 60°C for 3–5 min, washed in deionized water and air-dried.

The best metaphases were photographed using a 25 ISO Kodak imagilink film.

Quinacrine and chromomycin (CMA₃) staining

Chromosome staining with the AT-rich quinacrine fluorochrome followed the method described by Schmid (1980). The slides were first submitted to a gradual ethanol series (100%, 70% and 30%: 30 sec each) and immersed in a quinacrine solution (5 mg/100 ml) for 20 min at room temperature. Fluorescent signals for GC-rich regions were obtained using chromomycin A₃, according to Schweizer (1976). The slides were treated with 0.5 mg/ml CMA₃ for 1 h and counterstained with 0.1 mg/ml distamycin A (DA) for 15 min. The best metaphases were photographed using a 100 ISO Kodak T-MAX film, with an appropriated excitation filter.

Fluorescent *in situ* hybridization (FISH)

A fluorescent 18S rDNA probe was used for the localization of the ribosomal cistrons on the chromosomes. This was obtained from the nuclear DNA of *Salminus brasiliensis* (Pisces, Characidae), using the primers NS1 5'-GTAGTCATATGCTTGTCTC-3' and NS8 5'-TCCGCAG GTTACCTACGGA-3' (Hizume, 1994). After agarose gel analysis, the PCR products were purified with the Sephaglas Band Prep Kit (Pharmacia Biotech). Probes were labelled with biotinylated uridine (BdUTP), using the 'nick translation' kit, according to manufacturer instructions (Boehringer). Hybridization was detected with avidin-FITC and the chromosomes counterstained with propidium iodide. Chromosomal figures were obtained with a Kodak Gold Plus film, ISO 400, using a standard blue excitation filter.

Chromosomal analyses

About 30 metaphases were analyzed for each specimen and those of better quality were employed for the karyotype organization. At least six karyotypes were analyzed for each sex from each species/population. The arm ratio (Levan *et al.*, 1964) was used to classify the chromosomes according their morphology, through standardized measurements.

Results

Standard karyotype of *Triportheus*

All the *Triportheus* species/populations analyzed presented the diploid chromosome number $2n = 52$ for both sexes, and female heterogamety related to a ZZ/ZW sex chromosome system (Figures 1 and 2a, b). The Z chromosome corresponds to the biggest metacentric in the karyotype and the W to a medium-sized submetacentric chromosome.

C-banding

Almost all the chromosomes of the complement showed heterochromatin on the pericentromeric region. In the Z chromosome a clear C-band on both telomeric regions was also detected, while the W chromosome appeared strongly heterochromatic (Figure 1b).

NOR sites

The nucleolar organizer region (Ag-NOR), which is located on the short arm of a characteristic autosomal pair, showed a size polymorphism in all the species (Figures 1c and 2c (g-l)). This same chromosome presented C-positive bands on each side of the NOR site (Figures 1b and 2c (h)).

Sometimes additional Ag-NORs were found on the short arm of another unidentified autosome and on the long arm of the W chromosome. This occurs, for example, in *T. paranense* from the Cuiabá river (Figure 1c). In addition, a sporadic silver-stained region was also seen on the Z chromosome of *T. guentheri* (Figure 2c (c)).

Analysis of the NORs by FISH, using the 18S rDNA probe, confirmed the autosomal rDNA sites, as well as their location on the W chromosome (Figure 2b), in all the species/populations analyzed. These rDNA sites were the only chromosome regions showing positive and negative signals after chromomycin A₃ (Figure 2d) and quinacrine (Figure 2a) staining, respectively.

Discussion

Triportheus represents a genus of fish showing a constant diploid number ($2n = 52$) and a relatively conserved karyotypic macrostructure, as well as a differentiated ZZ/ZW sex chromosome system (Artoni *et al.*, 2001). While the Z is a conservative chromosome, corresponding to the biggest metacentric in the karyotype (Figure 1a), the W chromosome shows interspecific variations concerning its size, morphology and amounts of hetero- and euchromatin, being also largely heterochromatic (Figure 1b) and reduced in size in relation to the Z chromosome (Falcão, 1988; Artoni *et al.*, 2001; present study). However, some variation in the degree of heterochromatinization of the W chromosome can be found among the *Triportheus* species (Artoni *et al.*, 2001). The euchromatic regions of the W, although very reduced in some species, could represent chromosomal segments that still maintain a homology with the Z chromosome, as well as the physical locations of the genes for sex determination.

Base specific fluorochromes were not efficient in detecting different DNA classes in the *Triportheus* karyotype, with the exception of NORs. These regions appear as GC-rich according to the chromosome staining with chromomycin and quinacrine. Indeed, NORs have been

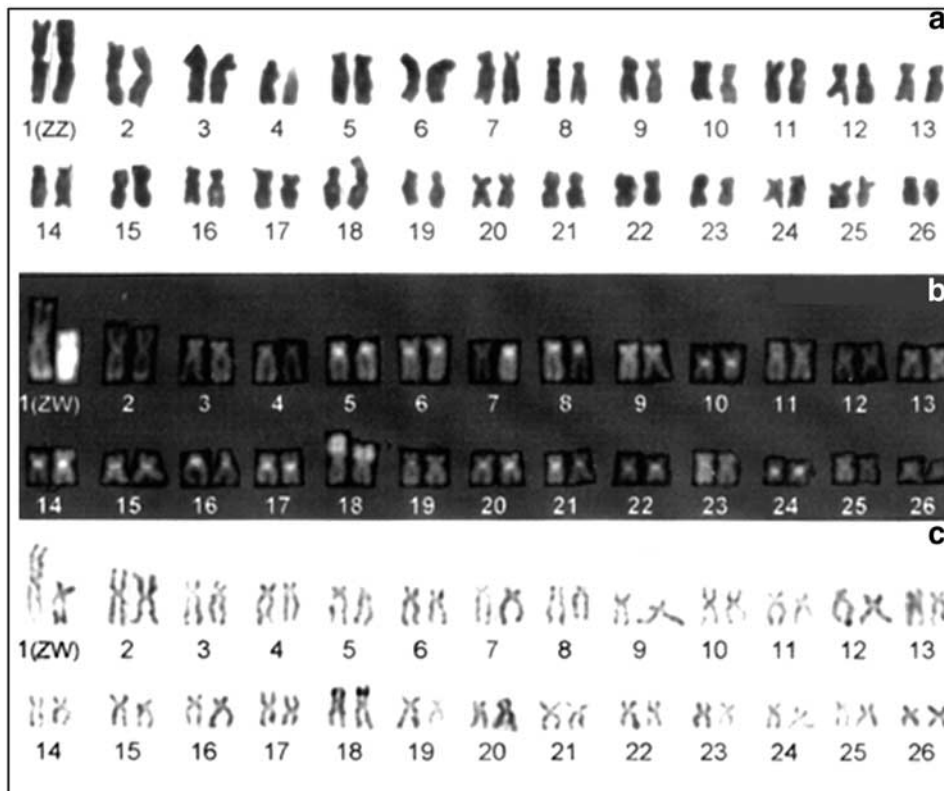


Figure 1 Male (a) and female (b, c) karyotypes of *Triportheus paranense* from the Cuiabá river, representatives for a general *Triportheus* karyotypic macrostructure. Conventional Giemsa staining (a), C-DAPI staining (b) and Ag-NORs (c).

characterized as GC-rich regions in lower vertebrates, such as amphibians (Schmid, 1980) and fishes (Amemyia and Gold, 1986; Phillips *et al.*, 1988; Galetti Jr and Rasch, 1993a, b; among others). However, GC-rich heterochromatic segments not related to the NORs can also be seen in some fish species (Artoni *et al.*, 1999).

A characteristic chromosome, bearing a secondary constriction on its short arm, appears to be a homeologous pair among the *Triportheus* species, showing a constant Ag-NOR (Figure 1). Sometimes another unidentified autosome pair was also stained with silver nitrate. Species with multiple NORs are not rare among fish (eg, Bertollo, 1996; Almeida-Toledo, 1998; Galetti Jr, 1998). In these cases numerical variation in Ag-NORs is frequently observed, which can be related to some factors, such as differential gene activation. The 'in situ' hybridization with a 18S rDNA probe was an important tool for the NORs studies in *Triportheus*. Thus, a maximum of four hybridization signals were observed in the autosomes, two of them not always detected due their small size. Additionally, a typical positive signal was always seen in the female specimens, on the long arm of the W chromosome (Figure 2b). This region is labelled by chromomycin A₃ (Figure 2d) and eventually is also stained with silver nitrate. No hybridization signal was detected in the Z chromosome, in spite of the sporadic occurrence of a silver-stained region on its long arm, as was seen in *T. guentheri* (Figure 2c (c)). This region may correspond to an argentophylic heterochromatin (Sumner, 1990), without any relationship to NORs, as observed in the *Hoplias lacerdae* fish group (Morelli, 1998).

Few examples of sex chromosomes bearing NORs are

known in mammals, especially those on the X chromosome, as in the bat *Carollia perspicillata* (Goodpasture and Bloom, 1975; Morielle and Varela-Garcia, 1988) and in the rodent *Akodon arviculoides* (Yonenaga-Yassuda *et al.*, 1983). These species do not show a significant difference in the number of active rRNA genes among males and females. A similar situation occurs in the amphibian *Gastrotheca riobambae*, where the females (XX) show two NORs and the males (XY) just one, without any evidence for a dosage compensation (Schmid *et al.*, 1983). Besides *Triportheus*, NORs located on fish sex chromosomes were already found in *Fundulus diaphanus* (Howell and Black, 1979), in the Arctic char *Salvelinus alpinus* (Reed and Phillips, 1997), and *Hoplias malabaricus* (Born and Bertollo, 2000).

Among invertebrates, such as *Drosophila* and many Coleoptera, where the synaptonemal complex does not occur between the X and Y chromosomes, the synapsis is assured by the association between the rDNA present in these chromosomes, allowing a regular segregation during gametogenesis (Irick, 1994; Ren *et al.*, 1997). Stitou *et al.* (1997) also described latent NORs in the sex chromosomes of the rodent *Lemniscomys barbarus*, suggesting some role for the rDNA during the synapsis process of the X and Y chromosomes. In this species, this kind of DNA was observed only in one population and a fast chromosomal evolution through translocations events (autosome/sex chromosomes) was proposed. However, in *Triportheus*, a peculiar condition can be characterized in view of the occurrence of inactive/active NORs on the W chromosome of the four species/populations analyzed. Thus, the location of rDNA sites on the sex chromosomes

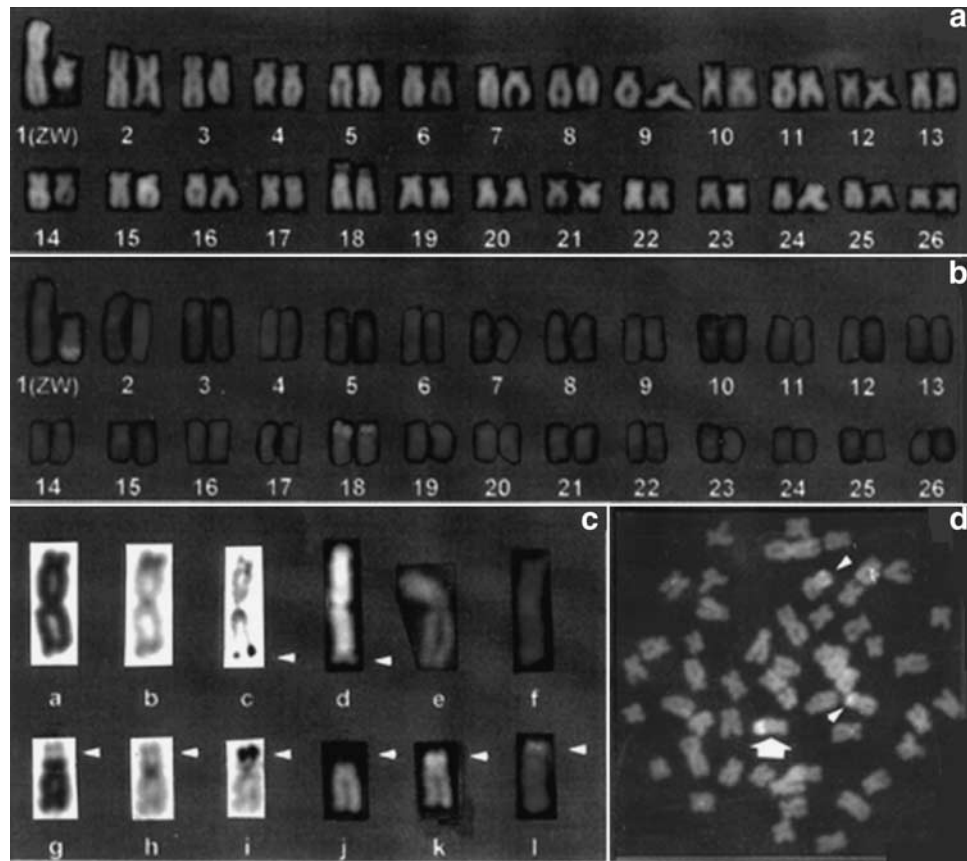


Figure 2 Karyotypes and chromosomes of *Triportheus paranense* from the Cuiabá river (a, b, d) and *T. guentheri* from the São Francisco river (c). Quinacrine staining (a) and fluorescence 'in situ' hybridization with 18S rDNA probe (b) showing differential signals on the long arm of the W chromosome and on the short arm of the 18th autosomal pair. Chromomycin-stained metaphase (d) showing labelled NORs in the 18th pair (arrowheads) and in the W chromosome (arrow). In (c) the Z chromosome (a–f) and the major NOR bearing autosome (g–l) of *T. guentheri* are shown, after conventional Giemsa staining (a,g), C-banding (b,h), silver staining (c,i), quinacrine staining (d,j), chromomycin A₃ staining (e,k) and FISH with 18S rDNA probe (f,l). The arrowheads indicate a sporadic silver-stained region on the Z chromosome and the NOR location on the autosome.

of *Triportheus* appears to be an old condition and probably already present in the ancestral W chromosome.

Some animal species offer special conditions for the study of sex chromosome differentiation. In the cassowary *Casuaris casuaris* (Aves, Ratitae), for instance, it was showed that the W chromosome is at an early stages of differentiation from the Z chromosome and that structural rearrangements might have been the initial step of this differentiation from an ancestral homomorphic pair. FISH analyses showed that some loci are conservative on both Z and W chromosomes, while others are missing from the W chromosome, probably due to deletions in its proximal region, leading to differences in the size and banding pattern between the Z and W chromosomes. Thus, these results favour previous propositions that heterochromatinization was not the initial step in the differentiation of the ratite W chromosome (Nishida-Umehara *et al*, 1999).

However, in several neotropical fish species, heterochromatinization is thought to be a plausible hypothesis concerning the evolution of the ZW sex chromosome system (Galetti Jr and Foresti, 1986; Moreira-Filho *et al*, 1993; Artoni *et al*, 2001). The same is true for the *Triportheus* species, in which the heterochromatinization might have driven the differentiation of the W chromosome and its

additional size reduction in respect to the Z chromosome (Artoni *et al*, 2001). In the Arctic char, *Salvelinus alpinus*, an important role was suggested for the rDNA loci on the putative sex chromosomes of this species, which might limit the opportunity for additional recombination near a major sex-determining locus, if this locus is adjacent to the rDNA sites (Reed and Phillips, 1997). In a similar way, we cannot discard a possible role for the repetitive DNA associated with the NORs, or even for the proper multicopy rDNA, in the differentiation of the sex chromosome system of *Triportheus*.

The ZZ/ZW sex chromosome system appears to be a synapomorphy in the *Triportheus* genus, representing a particular characteristic of this fish group among the large family Characidae. All of the nine species/populations already analyzed present a well differentiated ZZ/ZW system (Artoni *et al*, 2001). Accordingly, the occurrence of rDNA on the the W chromosome of this group may also correspond to a synapomorphic condition. Indeed, four different species/populations, from distinct hydrographic basins, present rDNA sites on an equivalent location in the W chromosome. Thus, further studies with other *Triportheus* species will offer additional support to the present proposition, as well as to the chromosomal evolution of this fish group.

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