

## Cytogenetic comparison of tree frogs of the genus *Aplastodiscus* and the *Hypsiboas faber* group (Anura, Hylidae)

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**ABSTRACT.** Four species of *Aplastodiscus* and two species of *Hypsiboas* were cytogenetically compared. *Aplastodiscus perviridis*, *A. cochranæ*, *H. albomarginatus*, and *H. faber* had  $2n = 24$  chromosomes, while *A. albosignatus* and *A. leucopygius* had  $2n = 20$  and  $2n = 18$  chromosomes, respectively. *Aplastodiscus perviridis* and *A. cochranæ* had identical karyotypes, as indicated by their chromosomal morphology, the location of the nucleolus organizer region (NOR) on chromosome pair 12, and the heterochromatin pattern. The NOR-bearing chromosomes of *A. albosignatus* and *A. leucopygius* (pair 9) were very similar in size and morphology (metacentric) when compared to *A. perviridis* and *A. cochranæ* (pair 12) and to *H. faber* (pair 11); the NOR of these chromosomes also occurred in the same region, suggesting that these chromosomes are homologous. Although *H. albomarginatus* differs from the other species with regard to the location of its NOR on pair 2, this species had the same diploid number and a chromosomal morphology similar to that of *A. perviridis* and *A. cochranæ*. Chromosomal differentiation among the species appears to have occurred by reduction in chromosome number.

**Key words:** *Aplastodiscus*; *Hypsiboas*; Karyotype; Nucleolus organizer region

## INTRODUCTION

The genus *Aplastodiscus* A. Lutz in B. Lutz (1950) currently consists of 15 species: *A. albofrenatus* (Lutz, 1924), *A. albosignatus* (Lutz and Lutz, 1938), *A. arildae* (Cruz and Peixoto, 1985), *A. calipygius* (Cruz and Peixoto, 1985), *A. cavicola* (Cruz and Peixoto, 1985), *A. cochranæ* (Mertens, 1952), *A. ehrhardti* (Muller, 1924), *A. eugenioi* (Carvalho e Silva and Carvalho e Silva, 2005), *A. flumineus* (Cruz and Peixoto, 1985), *A. ibirapitanga* (Cruz and Peixoto, 1985), *A. leucopygius* (Cruz and Peixoto, 1985), *A. musicus* (Lutz, 1949), *A. perviridis* (Lutz and Lutz, 1950), *A. sibilatus* (Cruz et al., 2003) and *A. weygoldti* (Cruz and Peixoto, 1987) (for review, see Faivovich et al., 2005; Frost, 2009). However, until the systematic revision of the Hylidae by Faivovich et al. (2005), this genus had only two species, *A. perviridis* A. Lutz in B. Lutz (1950) and *A. cochranæ* (Mertens, 1952). *Aplastodiscus perviridis* occurs from central and southeastern Brazil to northeastern Argentina (Cei and Roig, 1961; Frost, 2009), while *A. cochranæ* occurs at only three locations in the State of Santa Catarina, Brazil (Garcia et al., 2001).

The genus *Aplastodiscus* was initially separated from the genus *Hyla* Laurenti (1768) because of differences in finger and toe structure (A. Lutz in B. Lutz, 1950). However, since there are close morphological similarities between these two genera, some authors have doubted the validity of this separation, and subsequent publications continued to use the name *Hyla perviridis* instead of *A. perviridis* (Bokermann, 1967; Bokermann and Sazima, 1973; Cardoso et al., 1989; Cardoso and Haddad, 1992). The species *A. cochranæ* was described as *Hyla cochranæ* Mertens (1952), but Bokermann (1966) considered it synonymous to *Aplastodiscus perviridis*, a proposal accepted by various authors (Lutz, 1973; Duellman, 1977; Cei, 1980; Frost, 1985; Lavilla, 1992). Garcia et al. (2001) validated *A. cochranæ* based on external morphology.

According to Lutz (1950), the species *Hypsiboas albosignatus* (previously referred to as *Hyla albosignata*) is the closest to *A. perviridis* because of its size, color and song. Bokermann (1967) also reported similar song patterns for *A. perviridis*, *A. albofrenatus* (previously *Hyla albofrenata*) and *A. albosignatus* (previously *Hyla albosignata*). Another characteristic that approximates these species is the development of metacarpal and metatarsal tubercles, which could be related to the habit of digging burrows in mud (Garcia et al., 2001). A morphological phylogenetic analysis of the genus *Hyla* that included *Aplastodiscus* has corroborated the close relationship among these species (Silva, 1998).

Haddad et al. (2005) reported that *A. perviridis* had the same rare reproductive behavior, which is specific to species of the *A. albofrenatus* and *A. albosignatus* groups (referred to by Haddad as the *Hyla albofrenata* and *H. albosignata* complexes). Based on this similarity, these authors suggested a monophyletic origin for *Aplastodiscus* and these *Hyla* complexes.

Faivovich et al. (2005) provided a systematic review of the Hylidae based on molecular data. In this review, Brazilian species previously included in the “*albosignata*” and “*albofrenata*” complexes of the *Hyla albomarginata* group were transferred to the genus *Aplastodiscus*. In addition, the two species of the “*albomarginata*” complex of *Hyla*, e.g., *Hyla albomarginata*, were included in the *Hypsiboas faber* group. According to these authors, the genera *Aplastodiscus* and *Hypsiboas* are members of the tribe Cophomantini of Hyliinae.

There have been few cytogenetic studies of *Aplastodiscus* and of the species of the *Hypsiboas faber* group, with most of the karyotypes having been described using conventional techniques. Bogart (1973) found a distinct chromosome number for *A. albofrenatus* (Tijuca, RJ,  $2n = 24$  and Boracéia, SP,  $2n = 22$ ) and *A. albosignatus* (Boracéia, SP,  $2n = 20$  and Teresópolis, RJ,  $2n = 18$ ). In the *Hypsiboas faber* group, only the karyotype of *H. albomarginatus* ( $2n = 24$ ) (Beçak, 1968; Gruber et al., 2007) has been described.

In the present study, four species of *Aplastodiscus* representing the *A. perviridis* group (*A. perviridis* and *A. cochranae*) and the *A. albosignatus* group (*A. albosignatus* and *A. leucopygius*) and two species currently included in the *H. faber* group (*H. albomarginatus* and *H. faber*) were analyzed cytogenetically in order to assess the relationship between *A. perviridis* and *A. cochranae*, and between *Aplastodiscus* and the species of the *H. faber* group.

## MATERIAL AND METHODS

The collection sites and the number of specimens analyzed are shown in Table 1. Permissions for collecting the specimens were issued by the Instituto Brasileiro do Meio Ambiente e dos Recursos Naturais Renováveis (IBAMA - Proc. No. 02001.008867/01-92). All voucher specimens were deposited in the Célio Fernando Baptista Hadadd (CFBH) Collection, in the Department of Zoology of the State University of São Paulo (UNESP), Rio Claro, SP, Brazil.

**Table 1.** Data for the specimens analyzed.

| Species   | Locality<br>(municipality/state) | Number of<br>specimens analyzed | Voucher number<br>in the CFBH Collection |
|---|----------------------------------|---------------------------------|--|
| <i>Aplastodiscus perviridis</i> A. Lutz and B. Lutz, 1950   | Poços de Caldas (MG)             | 7 (6 m; 1 f)                    | 5840; 6987-6989; 7010-7012               |
|   | São Bento do Sul (SC)            | 5 (m)                           | 7406-7407; 5547-5549                     |
| <i>Aplastodiscus cochranae</i> Mertens, 1952                | Rancho Queimado (SC)             | 5 (m)                           | 6991; 7001; 7003-7005                    |
| <i>Aplastodiscus albosignatus</i> A. Lutz and B. Lutz, 1938 | Piraquara (PR)                   | 1 (m)                           | 5546                                     |
|   | São Bento do Sul (SC)            | 5 (m)                           | 5543-5545; 6992-6993                     |
| <i>Aplastodiscus leucopygius</i> Cruz and Peixoto, 1985     | Mogi das Cruzes (SP)             | 7 (m)                           | 4012-4013; 6646-6647; 7391-7393          |
|   | Maricá (RJ)                      | 2 (m)                           | 7395-7396                                |
| <i>Hypsiboas albomarginatus</i> Spix, 1824                  | Bertioga (SP)                    | 5 (m)                           | 6406-6410                                |
|   | Picinguaba (SP)                  | 4 (m)                           | 7408-7410; 4011                          |
|   | Mogi das Cruzes (SP)             | 2 (m)                           | 6403-6404                                |
| <i>Hypsiboas faber</i> Wied-Neuwied, 1821                   | Mogi das Cruzes (SP)             | 2 (m)                           | 4014; 6650                               |
|   | Biritiba Mirim (SP)              | 1 (m)                           | 6982                                     |

CFBH = Collection of the Department of Zoology of the State University of São Paulo (UNESP), Rio Claro, SP, Brazil. f = female, m = male.

Mitotic metaphases were obtained according to King and Rofe (1976) and Schimid (1978). The chromosomes were stained with 10% Giemsa in phosphate-buffered saline, pH 6.8. C-banding and Ag-NOR (nucleolus organizer region) techniques were carried out according to Sumner (1972) and Howell and Black (1980), respectively. Ag-NOR locations were confirmed with fluorescent *in situ* hybridization (Viégas-Péquignot, 1992) using a 28S rDNA probe of the recombinant plasmid HM123, which contains a fragment of rDNA of *Xenopus laevis* (Meunier-Rotival et al., 1979). The preparations were examined using an Olympus BX60 light microscope. The chromosomes were classified according to Green and Sessions (1991).

## RESULTS

All the specimens of *A. cochranae*, *A. perviridis*, *H. albomarginatus*, and *H. faber* had a diploid number of  $2n = 24$  chromosomes, whereas *A. albosignatus* had  $2n = 20$  and *A. leucopygius*,  $2n = 18$ . The chromosomal morphology was similar in all of these species, with pairs 1, 2, 9, 10, 11, and 12 being metacentric, and pairs 4, 5, 6, and 7 being submetacentric. There

was some variation in pair 3, which was metacentric in *H. albomarginatus* and submetacentric in the other species; similarly, pair 8 was submetacentric in *A. albosignatus* and *A. leucopygius* and metacentric in the other species (Figures 1A-F and 4A-F, Table 2). Heterochromatin was detected in *A. cochranae*, *A. perviridis* and *H. albomarginatus*. Heterochromatin blocks were observed in the centromeric regions of all chromosomes, but differed in amount. In *Aplastodiscus cochranae* and *A. perviridis*, heterochromatic blocks were detected in the telomeric region of the short arms of pairs 1 and 2 and in the long arms of pairs 1 to 8, as well as interstitially in the long arms of pairs 6, 10, 11, and 12, and in the short arms of pair 11 of both species. Pair 12 had heterochromatin adjacent to the NOR (Figures 2A-C and 4A, B,E). In *H. albomarginatus*, positive C-bands were found in the telomeric regions of the short arms of pairs 1, 2 and 3 and in the long arms of pairs 2 to 9. Interstitial heterochromatin was seen in the short arms of pairs 2 to 4 and in the long arms of pairs 5 and 6. On chromosome 2, the heterochromatin band coincided with the NOR and extended from this region to the centromere (Figures 2A-C and 4A,B,E).

The NOR occurred in the telomeric region of the long arm of pair 12 of *A. cochranae* and *A. perviridis* and of pair 9 of *A. albosignatus* and *A. leucopygius*. In *H. albomarginatus*, the NOR was located interstitially in the short arms of pair 2, and was heteromorphic for the homologous chromosomes. *Hypsiboas faber* had an NOR in the telomeric region of the long arm of pair 11 (Figures 3A-F and 4A-F). Secondary constrictions were frequently seen in some chromosomes of all species and were always associated with the NOR. In all species, the Ag-NOR coincided with the fluorescent *in situ* hybridization labeling obtained with the rDNA probe (Figure 3A-F). Table 3 summarizes the cytogenetic data for *Aplastodiscus* and the *Hypsiboas* species.

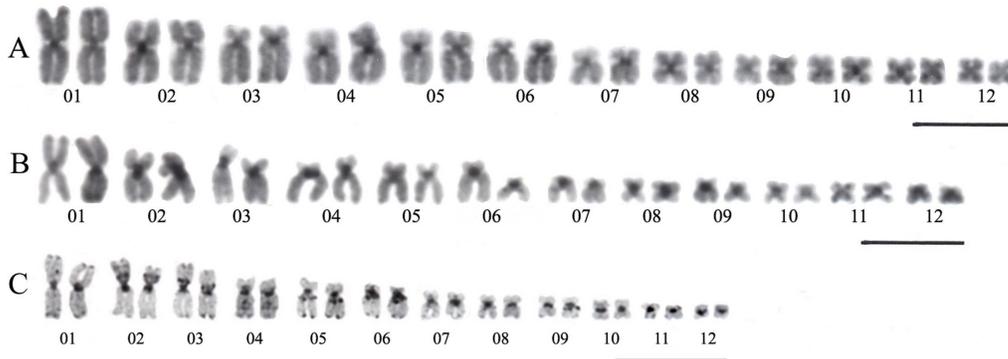
**Table 2.** Morphometrical data for chromosomes of *Aplastodiscus* and *Hypsiboas* species.

|                          | Chromosomes |                        |      |      |      |      |      |      |      |      |      |      |
|--------------------------|-------------|------------------------|------|------|------|------|------|------|------|------|------|------|
|                          | 1           | 2                      | 3    | 4    | 5    | 6    | 7    | 8    | 9    | 10   | 11   | 12   |
| <i>A. perviridis</i>     |             |                        |      |      |      |      |      |      |      |      |      |      |
| RL%                      | 14.9        | 12.7                   | 11.0 | 10.3 | 8.7  | 7.1  | 6.4  | 5.8  | 5.6  | 5.1  | 4.6  | 3.5  |
| CI                       | 0.45        | 0.41                   | 0.31 | 0.31 | 0.35 | 0.27 | 0.33 | 0.41 | 0.43 | 0.44 | 0.40 | 0.42 |
| CC                       | M           | M                      | SM   | SM   | SM   | SM   | SM   | M    | M    | M    | M    | M    |
| <i>A. cochranae</i>      |             |                        |      |      |      |      |      |      |      |      |      |      |
| RL%                      | 14.7        | 12.5                   | 10.6 | 10.0 | 8.8  | 7.2  | 6.5  | 5.8  | 5.6  | 5.3  | 4.9  | 4.8  |
| CI                       | 0.46        | 0.44                   | 0.27 | 0.35 | 0.34 | 0.30 | 0.37 | 0.45 | 0.45 | 0.46 | 0.45 | 0.48 |
| CC                       | M           | M                      | SM   | SM   | SM   | SM   | SM   | M    | M    | M    | M    | M    |
| <i>A. albosignatus</i>   |             |                        |      |      |      |      |      |      |      |      |      |      |
| RL%                      | 15.9        | 13.0                   | 11.3 | 10.9 | 10.3 | 10.0 | 8.7  | 6.6  | 4.9  | 2.2  |      |      |
| CI                       | 0.46        | 0.41                   | 0.33 | 0.31 | 0.29 | 0.29 | 0.29 | 0.27 | 0.40 | 0.45 |      |      |
| CC                       | M           | M                      | SM   | SM   | SM   | SM   | SM   | SM   | M    | M    |      |      |
| <i>A. leucopygius</i>    |             |                        |      |      |      |      |      |      |      |      |      |      |
| RL%                      | 17.9        | 13.9                   | 11.5 | 11.2 | 10.9 | 10.2 | 8.6  | 6.4  | 2.9  |      |      |      |
| CI                       | 0.48        | 0.39                   | 0.31 | 0.31 | 0.32 | 0.30 | 0.30 | 0.29 | 0.43 |      |      |      |
| CC                       | M           | M                      | SM   | SM   | SM   | SM   | SM   | SM   | M    |      |      |      |
| <i>H. albomarginatus</i> |             |                        |      |      |      |      |      |      |      |      |      |      |
| RL%                      | 15.3        | 15.1                   | 11.7 | 10.5 | 9.3  | 8.3  | 6.6  | 5.8  | 5.5  | 5.1  | 4.1  | 3.6  |
| CI                       | 0.48        | 13.2*<br>0.41<br>0.40* | 0.41 | 0.33 | 0.30 | 0.28 | 0.40 | 0.41 | 0.44 | 0.47 | 0.44 | 0.46 |
| CC                       | M           | M                      | M    | SM   | SM   | SM   | M    | M    | M    | M    | M    | M    |
| <i>H. faber</i>          |             |                        |      |      |      |      |      |      |      |      |      |      |
| RL%                      | 15.0        | 12.6                   | 10.9 | 10.3 | 8.8  | 7.1  | 6.4  | 5.7  | 5.5  | 5.1  | 4.6  | 4.0  |
| CI                       | 0.45        | 0.41                   | 0.30 | 0.34 | 0.35 | 0.28 | 0.32 | 0.46 | 0.42 | 0.44 | 0.40 | 0.44 |
| CC                       | M           | M                      | SM   | SM   | SM   | SM   | SM   | M    | M    | M    | M    | M    |

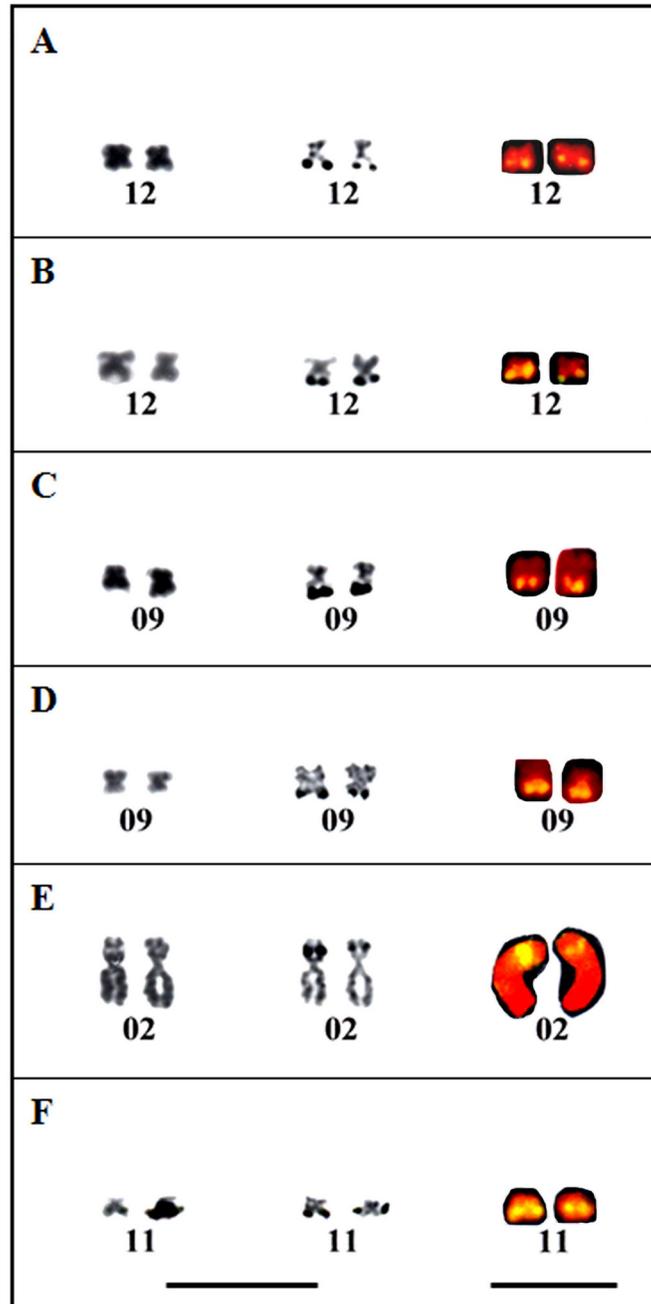
RL= relative length; CI= centromeric index; CC= centromeric classification; M= metacentric; SM= submetacentric. (\* = values for the morphs of chromosome 2 of *H. albomarginatus*).



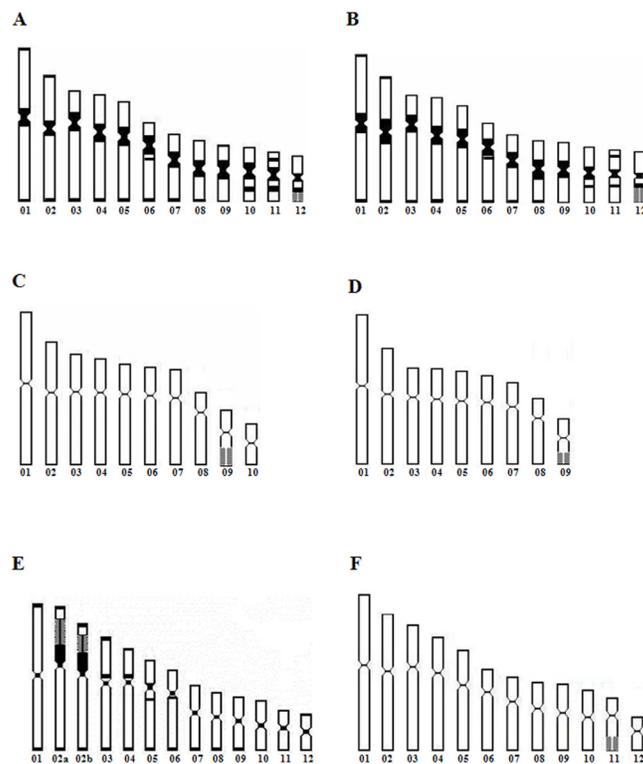
**Figure 1.** Karyotypes of *Aplastodiscus perviridis* (A), *A. cochranæ* (B), *A. albosignatus* (C), *A. leucopygius* (D), *Hypsiboas albomarginatus* (E), and *H. faber* (F). Arrows indicate the secondary constrictions. Bar = 10 µm.



**Figure 2.** Karyotypes of *Aplastodiscus perviridis* (A), *A. cochranæ* (B) and *Hypsiboas albomarginatus* (C) stained by C-banding. Arrows indicate interstitial and telomeric bands. Bar = 10 µm.



**Figure 3.** Chromosomal pairs containing the NOR region, as detected by Giemsa staining, silver staining and *in situ* hybridization with rDNA probe. *Aplastodiscus perviridis* (A), *A. cochranae* (B), *A. albosignatus* (C), *A. leucopygius* (D), *Hypsiboas albomarginatus* (E), and *H. faber* (F). Bar = 10  $\mu$ m.



**Figure 4.** Ideograms of the karyotypes of *Aplastodiscus perviridis* (A), *A. cochranae* (B), *A. albosignatus* (C), *A. leucopygius* (D), *Hypsiboas albomarginatus* (E), and *H. faber* (F). The black blocks indicate C bands and the gray areas show the NOR. The hatched regions are the secondary constrictions. In E, the letters *a* and *b* indicate the morphs of chromosome 2.

**Table 3.** Cytogenetic data for species of *Aplastodiscus* and *Hypsiboas*.

| Old taxon                       | Current taxon (sensu Faivovich et al., 2005) | 2n | Chromosomal morphology                     | NOR-bearing pair | Heterochromatin (non-centromeric C-bands) |                 |
|---------------------------------|--|----|--|------------------|---|-----------------|
|                                 |  |    |  |                  | Interstitial                              | Pericentromeric |
| <i>Aplastodiscus perviridis</i> | <i>Aplastodiscus perviridis</i>              | 24 | M; M; SM; SM; SM; SM; SM; M; M; M; M; M; M | 12q12q           | 11p, 11q and 10q                          | pair 6          |
| <i>Aplastodiscus cochranae</i>  | <i>Aplastodiscus cochranae</i>               | 24 | M; M; SM; SM; SM; SM; SM; M; M; M; M; M; M | 12q12q           | 11p and 10q                               | pair 6          |
| <i>Hyla albomarginata</i>       | <i>Hypsiboas albomarginatus</i>              | 24 | M; M; M; SM; SM; SM; M; M; M; M; M; M; M   | 2p2p             | 2p, 3p, 4p and 5 q, 6q                    | pair 2          |
| <i>Hyla faber</i>               | <i>Hypsiboas faber</i>                       | 24 | M; M; SM; SM; SM; SM; SM; M; M; M; M; M; M | 11q11q           | -   | -               |
| <i>Hyla albosignata</i>         | <i>Aplastodiscus albosignatus</i>            | 20 | M; M; SM; SM; SM; SM; SM; SM; M; M         | 9q9q             | -   | -               |
| <i>Hyla leucopygia</i>          | <i>Aplastodiscus leucopygius</i>             | 18 | M; M; SM; SM; SM; SM; SM; SM; M            | 9q9q             | -   | -               |

p = short arm; q = long arm.

## DISCUSSION

The analysis of chromosomal number and morphology has been useful for differentiating some frog species that have a similar external morphology, e.g., *Megaelosia* spp (Giretta and Aguiar-Jr, 1998; Rosa et al., 2003), *Dendropsophus nanus* and *D. sanborni* (as *Hyla* in Medeiros et al., 2003) and *Colostethus* (Veiga-Menoncello et al., 2003). As shown here, the karyotypes of *A. cochranae* and *A. perviridis* had the same diploid number of 24 chromosomes and a conserved chromosomal morphology. This characteristic is frequent in the Hylidae, as reported for closely related species such as *H. marginatus*, *H. semiguttatus* and *Hypsiboas* sp (aff. *H. semiguttatus*) (Ananias et al., 2004), *H. bischoffi* and *H. guentheri* (Raber et al., 2004), *H. polytenius* and *H. leptolineatus* (Vieira, 2004). However, in these species, the karyotypes could be distinguished by the NOR location and/or heterochromatin pattern. In contrast, in addition to having the same chromosomal number and morphology shown here, the NOR location and heterochromatin pattern were also identical in *A. cochranae* and *A. perviridis*, making it impossible to distinguish these two species based solely on their cytogenetic characteristics.

Although the two species of the *A. albosignatus* group, *A. albosignatus* and *A. leucopygius*, differed in their chromosomal number ( $2n = 20$  and  $18$ , respectively), their chromosomal morphology was very similar. The difference in chromosomal number apparently resulted from the loss of a small metacentric chromosome, probably chromosome 10, since chromosome pair 9 in the two species had almost the same size and had the NOR in the same position. Other chromosomes in these species were also morphologically very similar. The results for *A. albosignatus* differed from those reported by Bogart (1973), who analyzed two other populations assigned to this species, i.e., Boracéia, SP, with  $2n = 20$  and Teresópolis, RJ, with  $2n = 18$  chromosomes. The karyotype with  $2n = 18$  chromosomes found by Bogart is identical to that of *A. leucopygius* analyzed here and collected from a region (Maricá) very close (~50 km) to Teresópolis. Since *A. leucopygius* from the same locality (Teresópolis, RJ) was described by Cruz and Peixoto (1984), the cytogenetic data suggest that the species studied by Bogart (1973) was, in fact, *A. leucopygius* and not *A. albosignatus*.

The karyotypes of *A. albosignatus* ( $2n = 20$ ) and *A. leucopygius* ( $2n = 18$ ) were also similar to that of *A. cochranae* and *A. perviridis* ( $2n = 24$ ). The difference between these species appears to have resulted from a reduction in the number of chromosomes, since similar karyotypes, with  $2n = 24$ , occur in other Hylinae species. This reduction involves mainly the smaller chromosomes, since the first seven chromosomes are highly conserved in size and morphology. Also, the NOR-bearing chromosome 9 in *A. albosignatus* and *A. leucopygius* is very similar in size, morphology (metacentric) and NOR location (telomeric region) to pair 12 of *A. cochranae* and *A. perviridis*. A reduction in chromosome number has also been used to explain similar karyotypic variations in other anurans (Beçak, 1968; Bogart, 1970; Veiga-Menoncello et al., 2003; Siqueira-Jr. et al., 2004; Gruber et al., 2007). Although the chromosomal morphology of *H. albomarginatus* was very similar to that of the other species studied here, the location of the NOR on the large chromosome pair 2 differentiated this species from the others.

The NOR heteromorphism seen in the homologous chromosomes of pair 2 in *H. albomarginatus* is rather common in anurans and probably resulted from the amplification of rDNA sequences, which could explain the small increase in total length of the larger NOR-bearing chromosome.

The molecular data reported by Faivovich et al. (2005) showed that *H. albomarginatus* (previously *Hyla albomarginata*) was more closely related to other Hylinae species and was

included in the *H. faber* group, whereas *A. albosignatus* and *A. leucopygius* (previously *Hyla albosignata* and *Hyla leucopygia* of the *Hyla albomarginata* group) were related to *Aplastodiscus*. Both genera are in the same monophyletic tribe within the Hylinae. The karyotype and the NOR-bearing chromosomal pair in *H. faber* are very similar to those of *A. perviridis* and *A. cochranæ*, suggesting a close relationship, whereas in *H. albomarginatus*, the NOR location (pair 2) and the heterochromatin pattern are very distinct. The NOR in pair 11 of *H. faber* also occurs at the same site in *H. raniceps* (*H. albopunctata* group), whereas in *H. crepitans* (*H. faber* group) (Gruber et al., 2007) the NOR occurs at a site located interstitially. These findings suggest that the NOR sites and heterochromatin dispersion have been involved in the chromosomal evolution of this group.

The cytogenetic data obtained so far indicate that the species of *Aplastodiscus* and the *H. faber* group are very closely related and may have originated from a common ancestor. The chromosomal differentiation of the species analyzed here appears to have occurred by a reduction in the chromosome number (from  $2n = 24$  to  $2n = 20$  and  $2n = 18$ ), as already postulated by Gruber et al. (2007) to explain the reduction from  $2n = 24$  to  $2n = 22$  in *H. albopunctatus*, as well as by rearrangements involving mainly the smaller chromosomes (including the NOR-bearing ones). The latter conclusion is supported by the finding that the size and morphology of the first seven chromosomes are highly conserved in both genera.

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