

Comparative cytogenetics of some marsupial species (Didelphimorphia, Didelphidae) from the Amazon basin

Carlos Eduardo Faresin e Silva¹, Rodrigo Amaral de Andrade¹,
Érica Martinha Silva de Souza¹, Eduardo Schmidt Eler¹,
Maria Nazareth Ferreira da Silva², Eliana Feldberg¹

1 Laboratório de Genética Animal, Instituto Nacional de Pesquisas da Amazônia, Campus II, Avenida André Araújo, 2936, Manaus, Amazonas, Brazil **2** Coleção de Mamíferos, Instituto Nacional de Pesquisas da Amazônia, Campus II, Avenida André Araújo, 2936, Manaus, Amazonas, Brazil

Corresponding author: Carlos Eduardo Faresin e Silva (carlosfaresin@gmail.com)

Academic editor: T. Chassovnikarova | Received 3 June 2017 | Accepted 10 August 2017 | Published 26 October 2017

<http://zoobank.org/CAE22B3E-0F9F-4032-922A-71CEC447291B>

Citation: Silva CEF, Andrade RA, de Souza EMS, Eler ES, da Silva MNE, Feldberg E (2017) Comparative cytogenetics of some marsupial species (Didelphimorphia, Didelphidae) from the Amazon basin. *Comparative Cytogenetics* 11(4): 703–725. <https://doi.org/10.3897/CompCytogen.v11i4.13962>

Abstract

We investigated the karyotype of 18 didelphid species captured at 13 localities in the Brazilian Amazon, after conventional staining, C-banding, Ag-NOR and fluorescent *in situ* hybridization (FISH) using the 18S rDNA probe. Variations were found in the X chromosome, heterochromatin distribution and the 18S rDNA sequence. The main variation observed was in the position of the centromere in the X chromosome of *Caluromys philander* Linnaeus, 1758 and *Marmosa murina* Linnaeus, 1758. For both species, the X chromosome showed a geographical segregation in the pattern of variation between eastern and western Brazil, with a possible contact area in the central Amazon. C-banding on the X chromosome revealed two patterns for the species of *Marmosops* Matschie, 1916, apparently without geographic or specific relationships. The nucleolus organizer region (NOR) of all species was confirmed with the 18S rDNA probe, except on the Y chromosome of *Monodelphis touan* Shaw, 1800. The distribution of this marker varied only in the genus *Marmosa* Gray, 1821 [*M. murina* Thomas, 1905 and *M. demerarae* Thomas, 1905]. Considering that simple NORs are seen as a plesiomorphic character, we conclude that the species *Marmosa* spp. and *Didelphis marsupialis* Linnaeus, 1758 evolved independently to the multiple condition. By increasing the sample, using chromosomal banding, and FISH, we verified that marsupials present intra- and interspecific chromosomal variations, which suggests the occurrence of frequent chromosomal rearrangements in the evolution of this group. This observation contrasts with the chromosomal conservatism expected for didelphids.

Keywords

Marsupials, Amazon basin, C-band, NORs, 18S rDNA, Chromosomal rearrangements

Introduction

In the Americas, subclass Metatheria Huxley, 1880 is represented by the three marsupial orders: Didelphimorphia Gill, 1872, Paucituberculata Ameghino, 1894 and Microbiotheria Ameghino, 1889. The largest of the three American orders is Didelphimorphia, which is represented by the family Didelphidae Gray, 1821, whose species are widely distributed throughout the continent. Didelphidae is the only marsupial group present in Brazil. Together with rodents, they make up an important part of the mammalian fauna of the Amazon region (Voss and Jansa 2009, Wilson and Reeder 2011). Currently, 14 genera and 39 species are recorded in the Amazon basin. Although moderate in terms of species richness, didelphids are abundant in the region (Brandão et al. 2015).

Historically, the first cytogenetic data on American marsupials were recorded by Jordan (1911; cited in Reig et al. 1977), on the spermatogenesis of *Didelphis virginiana* Kerr, 1792. Since then, our knowledge of cytogenetics of American and Australian marsupials has grown significantly. Hayman (1990) reported the karyotype of 178 species of American and Australian marsupials and Svartman (2008) reported 45 karyotypes for American marsupials.

Unlike other mammal orders, such as Rodentia Bowdich, 1821, marsupials show relatively little chromosomal variation (Nagamachi et al. 2015). Chromosomal stability in marsupials was first verified in conventional staining karyotypes that revealed the existence of three main diploid numbers in species from both continents: 14, 18 and 22 chromosomes.

Among all the metatherian families, Macropodidae Gray, 1821 (order Diprotodontia Owen, 1866) is the most diverse in diploid number, varying from $2n=10$ to 32. While the American Didelphidae has only the three main diploid numbers, with the most frequent being $2n=14$ (Reig et al. 1977, Hayman 1990, Palma and Yates 1996, Carvalho et al. 2002), which has been suggested as the ancestral diploid number of all marsupials (Reig et al. 1977, Westerman et al. 2010). Further comparisons using chromosome banding in American and Australian marsupial species revealed that chromosomal stability is verified not only on the diploid number but also on longitudinal banding patterns that show intense conservation on chromatids (Yonenaga-Yassuda et al. 1982, Rofe and Hayman 1985, Casartelli et al. 1986, Souza et al. 1990, Svartman and Vianna-Morgante 1999).

Limited sampling effort has hampered the estimation of species richness in the Amazon, leaving large gaps in our knowledge of the mammalian fauna (Voss and Emons 1996, da Silva et al. 2001). Currently, of the 39 species of Amazonian marsupials (Brandão et al. 2015) only 17 have associated cytogenetic data (Nagamachi et al. 2015). However, considering the taxonomic instability of Amazonian marsupials,

this representation might not be accurate, since new phylogenetic studies will probably change the current classification of several taxa. Furthermore, the earlier literature often lacks a connection between the karyotype of putative species and the analyzed specimens, making it difficult to verify *a posteriori* the taxonomic identification attributed to a given karyotype.

The number of taxa analyzed to date is also limited, and existing cytogenetic analyses have been usually restricted only to the diploid and fundamental numbers (Nagamachi et al. 2015). New advances in the taxonomic classification of Amazonian marsupials, complementary techniques of cytogenetic analysis (banding, *in situ* hybridization), and added sampling efforts (more specimens, new localities) are necessary to improve current knowledge on the cytogenetics of these animals.

In this study, we analyze the main morphological differences in the sex chromosomes and the C-band pattern of 18 didelphid species from the Brazilian Amazon. In addition, we describe for the first time karyotype for six species (*Monodelphis touan*, *Monodelphis* aff. *adusta*, *Monodelphis* sp., *Marmosops impavidus*, *Marmosops bishopi* and *Marmosops pinheiroi*) and discuss these patterns in a broader geographical context, including other regions of Brazil and South America.

Material and methods

We cytogenetically analyzed 111 individuals in 18 species and 8 didelphid genera, collected in 13 localities in the Amazon (Table 1 and Figure 1). Scientific collecting permits were obtained from the Brazilian Institute of the Environment and Renewable Natural Resources (Instituto Brasileiro do Meio Ambiente e Recursos Naturais Renováveis – IBAMA), according to SISBIO license numbers: 02005.000642/03-11 (IBAMA); 02000.002336/2003-93 (IBAMA); 02005.002672/04 (IBAMA); 37585-5 (SISBIO); 37592-4 (SISBIO). The specimens were deposited at the Mammals Collection of the National Institute of Amazonian Research (INPA), Manaus, Brazil. Specimens are indicated by species, sampling sites, genus and collector number, followed by INPA collection number (in parentheses) when available, and their field codes are listed below. Karyotyped specimens at the figures: Figure 2: a) *Marmosa demerarae* (RNL 46, boxes: MCA 27); b) *Metachirus nudicaudatus* (SISTAP-M-302; boxes: SISIS-M-64); c) *Gracilinanus emiliae* (SISTAP-M-243); d) *Marmosa murina* (RNL 69, boxes: CEF 18); e) *Caluromys philander* (SISTAP-M-244, boxes: CAN 34, SISTAP-M-305); f) *Caluromys lanatus* (CTGA-M-701); ; g) *Marmosops pinheiroi* (INPA 5377, boxes: EE 192) (SISTAP-M-278, boxes: EE107, INPA 5408);. Figure 3: a) *Glironia venusta* (BAC 80); b) *Monodelphis* aff. *adusta* (INPA 5388); c) *Monodelphis touan* (INPA 5404); d) *Monodelphis* sp. (CAN 44); e) *Didelphis marsupialis* (EE 249, boxes: EE174)."

All voucher specimens: *Glironia venusta* Thomas, 1912: (BAC 80) – *Caluromys philander* Linnaeus, 1758: Tapajós River (male: SISTAP-M-297; SISTAP-M-305; SISTAP-M-318; SISTAP-M-382; female: SISTAP-M-244); Trombetas River (female:

Table 1. Didelphid species and their respective localities. Species analyzed in the current study were collected at localities 1 to 13, with number of individuals of males (M) and females (F) indicated. Geographic references for the current project were collected in a decimal degree projection using the WGS 84 reference. For literature data we insert converted geographical references where available. Localities with coordinates are presented only the first time they are cited in the table.

| Species | Locality | Locality Number | Coordinates† | M | F | Total | Reference |
|----------------------------|---------------------------------|-----------------|----------------------------------|---|----|---------------------|------------------------|
| <i>Caluromys philander</i> | Trombetas River, Pará State | 10 | 1.48163888889°S, 56.4573333333°W | 9 | 5 | 14 | Present work |
| | Tapajós River, Pará State | 11 | 3.35486111111°S, 55.2031666667°W | 1 | 1 | 2 | Present work |
| | Purus River, Amazonas State | 4 | 4.98066666667°S, 62.9770000000°W | | 1 | 1 | Present work |
| | Manaus, Amazonas State | 6 | 3.100548°S, 59.974595°W | | 1 | 1 | Present work |
| | Aragua, Venezuela | 14 | – | | | | Reig et al. 1977 |
| | Manaus, Amazonas State | 15 | 3.13333333333°S, 59.9500000000°W | | | | Souza et al. 2013 |
| | Jari, River, Pará State, Brazil | 12 | 0.7000000000°S, 52.6666666667°W | | 1 | 1 | Souza et al. 2013 |
| | Pernambuco state | 16 | – | | | | Souza et al. 1990 |
| São Paulo state | 17 | | | | | Pereira et al. 2008 | |
| <i>Caluromys lanatus</i> | Japurá River, Amazonas State | 1 | 1.84341666667°S, 69.0264722222°W | | 1 | | Present work |
| | Iquitos, Peru | – | – | | | | Hayman and Martin 1974 |
| | Manaus, Amazonas State | – | – | | | | Casartelli et al. 1986 |
| | Rorôndia, Brasil | 13 | – | | | | Souza et al. 1990 |
| <i>Marmosa demerarae</i> | Aripuanã River, Amazonas State | 7 | 6.0000000000°S, 60.1666666667°W | 4 | 4 | 8 | Present work |
| | Manaus, Amazonas State | 6 | 3.13333333333°S, 59.9500000000°W | 7 | 11 | 18 | Present work |
| | Cuieiras River, Amazonas State | 5 | 2.70708611111°S, 60.3738388889°W | 4 | 2 | 6 | Present work |
| | Purus River, Amazonas State | 4 | 0.5772500000°S, 64.8976944444°W | 3 | 4 | 7 | Present work |
| | Negro River, Amazonas State | 3 | 0.5772500000°S, 64.8976944444°W | 1 | 5 | 7 | Present work |
| | Tapajós River, Pará State | 11 | 3.35486111111°S, 55.2031666667°W | 3 | 5 | 9 | Present work |
| | Trombetas River, Pará State | 10 | 1.48163888889°S, 56.4573333333°W | 9 | 5 | 14 | Present work |
| | Jari River, Pará State | 12 | 0.7000000000°S, 52.6666666667°W | 9 | 2 | 11 | Present work |
| | Juruá River, Amazonas State | 2 | 3.64151111111°S, 66.1006916667°W | | 1 | 1 | Present work |
| | Jatapú River, Amazonas State | 9 | 2.017940°S, 58.203228°W | | 1 | 1 | Present work |
| | Jari River, Pará State | 12 | 0.7000000000°S, 52.6666666667°W | 1 | | 1 | Present work |
| | Uatumá River, Amazonas State | 8 | 1.84998888889°S, 59.4402000000°W | 5 | 3 | 9 | Present work |
| | Trombetas River, Pará sate | 10 | 1.48163888889°S, 56.4573333333°W | | 1 | 1 | Present work |
| | Negro River, Amazonas State | 3 | 0.5772500000°S, 64.8976944444°W | 1 | 1 | 2 | Present work |
| | Juruá River | 2 | 3.64151111111°S, 66.1006916667°W | | 1 | 1 | Present work |

| Species | Locality | Locality Number | Coordinates† | M | F | Total | Reference |
|--|--|------------------|-----------------------------------|---|---|----------------------|------------------------|
| <i>Marmosa murina</i> | Purus River, Amazonas State | 4 | 0.57725000000°S, 64.89769444444°W | 2 | | 2 | Present work |
| | Pernambuco State | 16 | – | | | | Souza et al. 1990 |
| | Villa Vivencio, Colombia | 18 | – | | | | Hayman and Martin 1974 |
| | Bolivar, Venezuela | 19 | – | | | | Reig et al. 1977 |
| | Tartarugalzinho, Amapá State | 21 | | | | | Carvalho et al. 2002 |
| | Loreto, Peru | 20 | – | | | | Reig et al. 1977 |
| | Vila Rica, Mata Grosso State | 22 | 10°01'S, 51°07'W | | | | Pagnozzi et al. 2002 |
| | UHE Peixe Angical, Tocantins State | 23 | 12°01'30"S, 48°32'21"W | | | | Pereira et al. 2008 |
| | Porto Nacional, Tocantins state | 24 | 10°42'S, 48°25'W | | | | Lima 2004 |
| | Uruaçu, Goiás state | 25 | 14°31'S, 49°08'W | | | | Carvalho et al. 2002 |
| | Colinas do Sul, Goiás state | 26 | 14°09'S, 48°04'W | | | | Carvalho et al. 2002 |
| | UHE Corumbá IV Luziania, Goiás state | 27 | 16°15'09"S, 47°57'01"W | | | | Carvalho et al. 2002 |
| | Pacoti, Ceará state | 28 | 4°13'S, 38°55'W | | | | Pagnozzi et al. 2002 |
| Reserva Biológica Duas Bocas, Espírito Santo state | 29 | 20°16'S, 40°28'W | | | | Paresque et al. 2004 | |
| <i>Gracilinanus emiliae</i> | Tapajós River, Pará state | 11 | 35486111111°S, 55.2031666667°W | 3 | 1 | 4 | Present work |
| | Serra da Mesa, Colinas do Sul, Goiás state | 26 | 14°09'S 48°04'W | | | | Carvalho et al. 2002 |
| | UHE Corumbá IV, Luziania, | 27 | 16°15'09"S, 47°57'01"W | | | | Pereira et al. 2008 |
| <i>Metachirus nudicaudatus</i> | Trombetas River, Pará state | 10 | 1.48163888889°S, 56.45733333333°W | | 1 | 1 | Present work |
| | Jari River, Pará state | 12 | 0.7000000000°S, 52.6666666667°W | 1 | | 1 | Present work |
| | Cuieiras River, Amazonas state | 5 | 2.70708611111°S, 60.3738388889°W | 1 | | 1 | Present work |
| | Juruá River, Amazonas state | 2 | 3.64151111111°S, 66.1006916667°W | | 1 | 1 | Present work |
| | Tapajós River, Pará state | 11 | 3.54861111111°S, 55.2031666667°W | 2 | 2 | 4 | Present work |
| <i>Glironia venusta</i> | Porto Velho, Rondônia State | 13 | 8.87416666667°S, 64.0077777778°W | | 1 | 1 | Present work |
| <i>Monodelphis touan</i> | Jari River, Pará state | 12 | 0.7000000000°S, 52.6666666667°W | 3 | | 3 | Present work |
| <i>Monodelphis</i> sp. | Purus River, Amazonas state | 4 | 0.57725000000°S, 64.89769444444°W | | 1 | 1 | Present work |
| <i>Monodelphis aff. adusta</i> | Aripuanã River, Amazonas state | 7 | 6.00000000000°S, 60.1666666667°W | 1 | | 1 | Present work |
| <i>Monodelphis emiliae</i> | Aripuanã River, Amazonas state | 7 | 6.00000000000°S, 60.1666666667°W | | 1 | 1 | Present work |
| | Juruá River, Acre state | | 8°40'S 72°47'W | | | | Patton et al. 2000 |
| <i>Monodelphis breviceaudata</i> | Negro River state | 3 | 0.57725000000°S, 64.89769444444°W | 1 | | 1 | Present work |
| <i>Marmosops bishop</i> | Aripuanã River, Amazonas state | 7 | 6.00000000000°S, 60.1666666667°W | 5 | 6 | 11 | Present work |
| | Purus River, Amazonas state | 4 | 0.57725000000°S, 64.89769444444°W | 2 | 1 | 3 | Present work |
| | Negro River, Amazonas state | 3 | 0.57725000000°S, 64.89769444444°W | 1 | | 1 | Present work |

| Species | Locality | Locality Number | Coordinates† | M | F | Total | Reference |
|------------------------------|--|-----------------|----------------------------------|---|---|-------|----------------------|
| <i>Marmosops pinheiroi</i> | Tapajós River, Pará state | 11 | 3.54861111111°S, 55.2031666667°W | 4 | 2 | 6 | Present work |
| | Trombetas River, Pará state | 10 | 1.48163888889°S, 56.4573333333°W | 8 | 1 | 9 | Present work |
| | Cuieiras River, Amazonas state | 5 | 2.70708611111°S, 60.3738388889°W | 3 | 2 | 5 | Present work |
| <i>Marmosops parvidens</i> | Jari River, Pará state | 12 | 0.7000000000°S, 52.6666666667°W | | 2 | 2 | Present work |
| | Jatapú River, Amazonas state | 9 | 2.017940°S, 58.203228°W | 4 | 3 | 7 | Present work |
| | La Paz, Bolívia | – | – | | | | Palma and Yates 1996 |
| | Serra da Mesa, Colinas do Sul, Goiás state | 26 | 14°09'S, 48°04'W | | | | Carvalho et al. 2002 |
| | Apiacás, Mato Grosso state | | 9°34'S, 57°23'W | | | | Pagnozzi et al. 2002 |
| <i>Marmosops impavidus</i> | Juruá River, Amazonas state | 2 | 3.64151111111°S, 66.1006916667°W | 2 | 1 | 3 | Present work |
| <i>Marmosops pakaraimae</i> | Japurá River, Amazonas state | 1 | 1.84341666667°S, 69.0264722222°W | | 1 | 3 | Present work |
| <i>Didelphis marsupialis</i> | Tapajós River, Pará state | 11 | 3.54861111111°S, 55.2031666667°W | 1 | 3 | 4 | Present work |
| | Trombetas River, Pará state | 10 | 1.48163888889°S, 56.4573333333°W | 1 | 2 | 3 | Present work |
| | Manaus, Amazonas state | 6 | 3.13333333333°S, 59.9500000000°W | 8 | 4 | 12 | Present work |
| | Uatumá River, Amazonas state | 9 | 2.017940°S, 58.203228°W | 1 | 1 | 2 | Present work |
| | Cuieiras River, Amazonas state | 5 | 2.70708611111°S, 60.3738388889°W | 2 | 2 | 4 | Present work |

CTGA-M-652); Purus River (female: CAN 34); Manaus (female: MSN 01); (female: BAC 102) – *Caluromys lanatus* Olfers, 1818: Japurá River (female: CTGA-M-701) – *Marmosops* sp. Matschie, 1916: Aripuanã River (female: MCA 3; MCA 7; MCA 8; MCA 26; MCA 31; MCA 35; male: MCA 4; MCA 16; MCA 38; MCA 39); Jari River (female: TAG 3459; RNL 70); Juruá River (male: EE 107; EE 139; female: EE135); Cuieiras River (female: EE 198; EE 211; male: EE 192; EE 201; EE216) – *Marmosops bishopi* Pine, 1981: Negro River (male: SISIS-M-127); Purus River (male: SISPUR-M-135; SISPUR-M-157; SISPUR-M-160; SISPUR-M-164; SISPUR-M-135; CAN 30; CAN 51; female: CAN 48) – *Marmosops pinheiroi* Pine, 1981: Tapajós River (male: SISTAP-M-237; SISTAP-M-278; female: SISTAP-M-268; SISTAP-M-277) – *Marmosops parvidens* Tate, 1931: Trombetas River (male: CTGA-M-501; CTGA-M-516; CTGA-M-531; CTGA-M-532; CTGA-M-551; CTGA-M-555; CTGA-M-581; CTGA-M-600; female: CTGA-M-533) – *Marmosops impavidus* Tschudi, 1845: Purus River (male: SISPUR-M-149) – *Marmosops* cf. *pakaraimae* Voss, Lim, Díaz-Nieto et Jansa 2013: Japurá River (male: SISJAP-M-705) – *Marmosa murina* Linnaeus, 1758: Jari River (male: RNL 45); Uatumá River (male: CEF 4; CEF 8; CEF 18; CEF 27; CEF 28; CEF 32; female: CEF 16; CEF 34; CTGA-M-8; CTGA-M-22; CTGA-M-41;), Negro River (male: SISIS-M-57; SISIS-M-63); Trombetas River (female: CTGA-M—519); Purus River (male: CAN 43); Japurá River (male: CTGA-M-708)

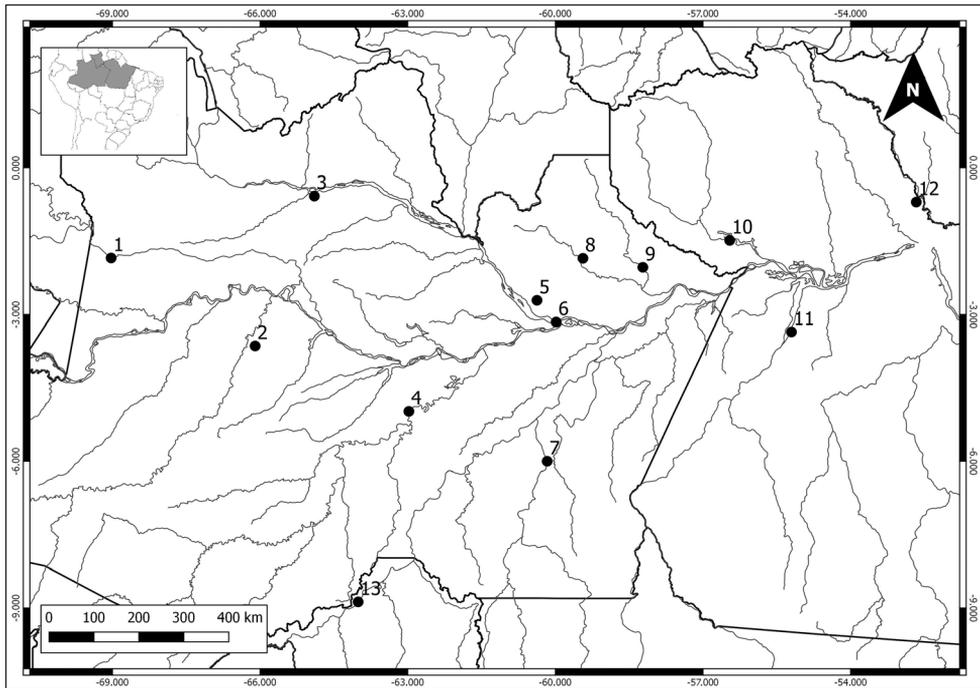


Figure 1. Sampling sites plotted on the Amazon basin map, Amazonas State: **1** Japurá River, Japurá city **2** Juruá River, Juruá city **3** Negro River, Santa Isabel do Rio Negro city **4** Purus River, Tapauá city **5** Cuieiras River, Manaus city **6** Manaus city, urban área: Federal University of Amazonas's campus (UFAM) and Isaac Sabá Oil Refinery **7** Aripuaná River, Novo Aripuaná city **8** Uatumá River, Presidente Figueiredo city **9** Jatapú River, São Sebastião do Uatumá city; Pará State: **10** Trombetas River, Oriximiná city **11** Tapajós River, Aveiro and Santarém cities **12** Jari River, Almeirim city; Rondônia State: **13** Madeira River, Porto Velho city. Geographic coordinates at the Table 1.

– *Marmosa murina* Linnaeus, 1758: Aripuaná River (female: MCA12, Japurá River (male: SISJAP-M-764)- *Gracilinanus emiliae* Thomas, 1909: Tapajós River: (male: SISTAP- M-245; SISTAP- M-343; SISTAP- M-344; SISTAP- M-345) – *Marmosa demerarae* Thomas, 1905: Aripuaná River (female: MCA 27; MCA 36; MCA 58; MCA 65; male: MCA 21; MCA 59); Jari River (female: RNL 31; RNL 48; male: RNL 30; MCA 32; MCA 46; MCA 49; MCA 58; MCA 61; MCA 64; MCA 66; MCA 67) Juruá River (female: EE136; male: EE 143); Manaus (female: EE 149; EE 150; EE 151; EE 154; EE 158; EE 159; EE 169; EE 222; EE 228; 229; EE 234; male: EE 157; EE 167; EE 170; EE 176; EE 189; EE 194; EE 196; EE 202; EE 215; EE 220; EE 235); Cuieiras River (female: EE 193; EE 219); Tapajós River (female: SISTAP-M-229; SISTAP-M-241; SISTAP-M-321; SISTAP-M-333; SISTAP-M-369; male: SISTAP-M-267; SISTAP-M-279; SISTAP-M-322); Trombetas River (female: CTGA-M-579; CTGA-M-590; CTGA-M-622; CTGA-M-667; CTGA-M-672; male: CTGA-M-535; CTGA-M-539; CTGA-M-557; CTGA-M-558; CTGA-M-572; CTGA-M-573; CTGA-M-578; CTGA-M-580; CTGA-M-613); Negro River (female:

SISIS-M-85; SISIS-M-110; SISIS-M-117; SISIS-M-128; male SISIS-M- 86); Purus River (female: SISPUR-M-145; CAN 25; CAN 31; CAN 50: male: SISPUR-M-144; SISPUR-M-147; SISPUR-M-148) – *Monodelphis* aff. *adusta* Thomas, 1897: Madeira River (male: MCA 15) – *Monodelphis touan*: Jari River (male: TAG 2731; RNL 68) – *Monodelphis* sp. Burnett, 1830: Purus River: (male: CAN 44) – *Monodelphis emiliae* Thomas, 1912: Aripuaná River (female: MCA 31) – *Metachirus nudicaudatus* Geoffroy et Saint-Hilaire, 1803: Jari: River (RNL 47); Cuieiras River: (female: EE 200); Tapajós River (female: SISTAP-M-230; SISTAP-M-230; male: SISTAP-M-251; SISTAP-M-269); Trombetas River: (female: CTGA-M-655); Jatapú River: (female: CTGA-M-52; CTGA-M-58); Negro River: (female: SISIS-M-64; SISIS-M-78; male: SISIS-M-84; SISIS-M-116); Purus River: (male: CAN 33) – *Didelphis marsupialis* Linnaeus 1758: Jari River: (female: RNL 44; RNL 53; RNL 59; male: RNL 52; RNL 55; RNL 62; RNL 63); Manaus: (female EE 174; EE 197; EE 204; EE 224; EE 227; EE 246; EE 250; EE 155; EE 155; EE 173; EE 183; EE 190; EE 203; EE 205; EE 206; EE 223; EE 232; EE 233; EE 237; EE 247; EE 248; EE249; EE 190); Uatumã River (female: CEF 5; male: CEF 13); Trombetas River (female: CTGA-M-594; CTGA-M-606; male: CTGA-M-607); Purus River (male: SISPUR-M-185); Negro River (male: SISIS-M-73); Tapajós River (female: SISTAP-M-324; SISTAP-M-346; SISTAP-M-347; male: SISTAP-M-243); Japurá River: (male: CTGA-M-732).

The metaphases were obtained from bone marrow by *in vivo* method according to Ford and Harmerton (1956). Each animal received 1mL/100g weight of a 0,0125% colchicine solution for 30 minutes, the cells were exposed for 20 minutes to a 0,075M KCl solution, fixed 3:1 in methanol and acetic acid and stored at -20 °C. The C-band and Nucleolus Organizing Regions (NORs) patterns were determined according to the techniques described by Sumner (1972), and Howell and Black (1980), respectively. Chromosome pairing considered morphology in decreasing order of size and the chromosomes were classified as metacentric (m), submetacentric (sm), subtelocentric (st) and acrocentric (a) according to the ratio of chromosome arms and the position of the centromere, according to Patton (1967). 18S rDNA sequences were mapped by fluorescence *in situ* hybridization (FISH) according to Pinkel et al. (1986), whose probe was obtained by polymerase chain reaction (PCR) using the following primers designed by Gross et al. (2010): 18SF (5'-CCG CTT TGG TGA CTC TTG AT-3') e 18SR (5'-CCG AGG ACC TCA CTA AAC CA-3') and labeled with digoxigenin (DIG-Nick translation, ROCHE) or Biotin (Bio-Nick translation, ROCHE), following manufacturer's instructions.

Results

Among the 18 species analyzed, 11 showed $2n=14$; six $2n=18$ and one $2n=22$ chromosomes (Table 1).

In the species with $2n=14$, we observed a very similar structure among the autosomes. These karyotypes include six autosome pairs (Fig. 2), three large submetacentric

pairs, one metacentric pair and two small pairs that varied in morphology in the different species, resulting in differences in the chromosomal formulas and fundamental numbers (FNa). FNa=20, with formula $2m+6sm+4a+XX/XY$, was recorded in *Marmosa demerarae* (Fig. 2a-I) and *Metachirus nudicaudatus* Geoffroy an Saint-Hilaire, 1803 (Fig. 2b-I). FNa=22, with formula $2m+6sm+2st+2a+XX/XY$, was present in *Gracilinanus emiliae* Thomas, 1909 (Fig. 2c-I), *Marmosa murina* (Fig. 2d-I), *Caluromys philander* (Fig. 2e-I) and *Caluromys lanatus* Olfers, 1818 (Fig. 2f-I). FNa=24, with formula $6m+6sm+XX/XY$, was recorded in species of the genus *Marmosops* including *M. bishopi* (Pine, 1981), *M. pinheiroi* Pine, 1981, *M. parvidens* Tate, 1931, *M. impavidus* Tschudi, 1845, and *M. cf. pakaraimae* Voss, Lim, Díaz-Nieto et Jansa 2013. The five species of *Marmosops* presented similar karyotypic characteristics (Fig. 2g-I – only *M. pinheiroi* shown).

We observed three different morphologies for X chromosome: metacentric in *G. emiliae* and *Marmosops* spp. (Fig. 2c-I and 2g-I); submetacentric in the only female of *C. lanatus* (Fig. 2f-I); and acrocentric in *M. demerarae* and *M. nudicaudatus* (Fig. 2a-I and 2b-I). In *Caluromys philander* and *Marmosa murina*, we observed an intraspecific variation in the structure of the X chromosome, acrocentric and submetacentric, both in specimens from the same and different localities (Fig. 2e-I and 2d-I).

The bare-tailed woolly opossum (*C. philander*) has X chromosome either acrocentric or submetacentric, with females being either homozygous or heterozygous carriers of the heteromorphic X (Fig. 4). In the murine mouse opossum (*Marmosa murina*), the metacentric or submetacentric X was found in individuals throughout the Brazilian Amazon, except in the Purus River (Fig. 5, locality 4); it was also found in individuals from two localities in central Brazil (Fig. 5, localities 25 and 26). These are situated at the southern limits of the distribution of *M. murina* and both, the submetacentric X and the acrocentric X, are sympatric at locality 26. Furthermore, in the northern Amazon in Colombia, Venezuela and Peru, the X chromosome is metacentric (Fig. 5, localities 18, 19 and 20) (Reig et al. 1977, Carvalho et al. 2002). The acrocentric X was found in the Purus River (Fig. 5, locality 4), and in central, southeastern and northeastern Brazil (Fig. 5, localities 16 and 22-28) (Souza et al. 1990, Palma and Yates 1996, Carvalho et al. 2002).

The Y chromosome was acrocentric in *G. emiliae*, *Marmosops* spp., *M. demerarae* and *M. nudicaudatus* (Fig. 2c, g, a, b), and dot-like in *C. philander* and *M. murina* (Fig. 2e, d).

Among the species with $2n=18$ chromosomes, FNa=20 was recorded in *Glironia venusta* Thomas, 1912, with formula $2m+2sm+2st+10a+XX/XY$ (Fig. 3a-I), FNa=30 was recorded in four species of the genus *Monodelphis* Burnett, 1830: *M. aff. adusta* Thomas, 1897 (Fig. 3b-I), *M. touan* (Fig. 3c-I), *M. emiliae* Thomas, 1912 (Fig. 3d-I), and *M. brevicaudata* Erxleben, 1777 (Fig. 3e-I) with formula $2m+2sm+8st+2a+XX/XY$, and FNa=32 in *Monodelphis* sp. (Fig. 3f-I), with formula $2m+2sm+10st+2a+XX/XY$. We observed two X chromosomes morphologies: acrocentric in *M. aff. adusta*, *M. touan* and *M. brevicaudata* (Fig. 3b, c, e), and submetacentric in *Monodelphis emiliae* and *Monodelphis* sp. (Fig. 3d, f). The Y chromosome was acrocentric in *M. touan* and *Monodelphis* sp., and dot-like in *M. aff. adusta* and *M. brevicaudata*.

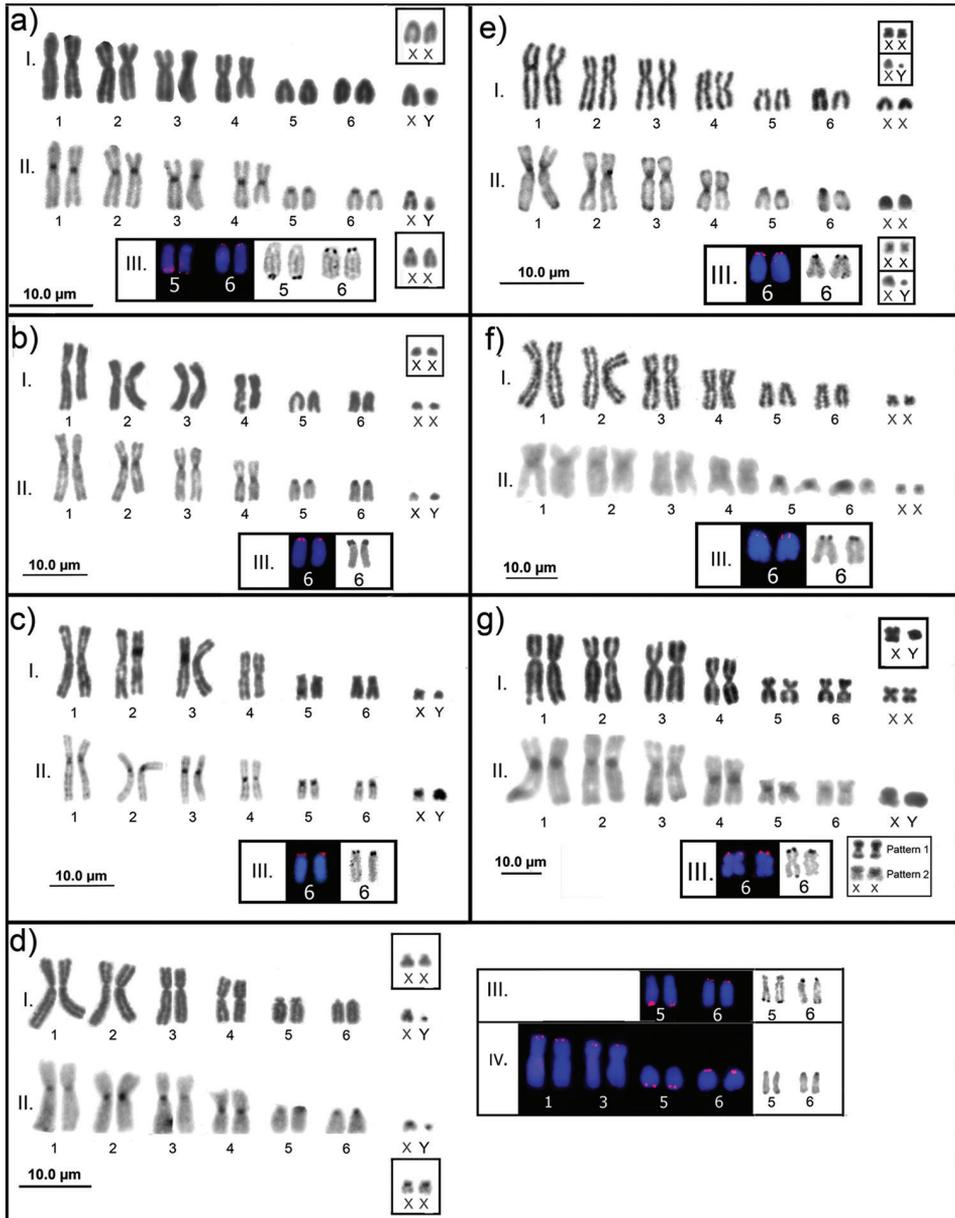


Figure 2. Karyotypes under conventional staining (I), C-band (II), 18S rDNA and Ag-NOR (III), sex chromosomes in the boxes: **a** *Marmosa demerarae* **b** *Metachirus nudicaudatus* **c** *Gracilinanus emiliae* **d** *Marmosa murina*, (IV) variations on the 18S sites found in the individuals from Purus River, Tapauá city, Amazonas State **e** *Caluromys philander* **f** *Caluromys lanatus* **g** *Marmosops pinheiroi*.

Didelphis marsupialis was the only species that presented $2n=22$ chromosomes and $FN_a=20$, with formula $20a+XX/XY$ (Fig. 3g-I), with acrocentric X and Y.

The position of the heterochromatin on the $2n=14$ species was centromeric, being conspicuous in *M. demerarae* (Fig. 2a-II), *M. nudicaudatus* (Fig. 2b-II), *G. emiliae*

(Fig. 2c-II), *M. murina* (Fig. 2d-II), and *M. pinheiroi* (Fig. 2g-II). *Caluromys philander* and *C. lanatus* exhibited tenuous heterochromatin, with additional telomeric heterochromatin in *C. philander* chromosomes (Fig. 2e-II and 2f-II). The X chromosome in *C. philander* was entirely heterochromatic, except for a distal band in the long arms (Fig. 2e-II); in *M. demerarae* it was also entirely heterochromatic, except for a proximal euchromatic band in the long arms (Fig. 2a-II); in *M. murina* (Fig. 2d-II), *M. nudicaudatus* (Fig. 2b-II) and *G. emiliae* the heterochromatin was centromeric (Fig. 2c-II).

Two C-band patterns were present in the X chromosome for species of *Marmosops*. In pattern 1, X was entirely heterochromatic except for a proximal band in the long arms (Fig. 2g – box); in pattern 2, the heterochromatin was in the short arms and the centromere (Fig. 2g – box). Both patterns were present in *M. parvidens* and *M. bishopi*, while only pattern 1 was observed in *M. cf. pakaraimae*, *M. impavidus* and *M. pinheiroi* (Table 2). The Y chromosome was entirely heterochromatic in all species.

In the species with $2n=18$ chromosomes, the heterochromatin was centromeric in *G. venusta* (Fig. 3a-II), *M. aff. adusta* (Fig. 3b-II), *M. touan* (Fig. 3c-II) and *M. emiliae* (Fig. 3d-II). The Y chromosome was entirely heterochromatic in *M. adusta* (Fig. 3b-II) and *M. touan* (3c-II). It was not possible to determine the C-band pattern in *Monodelphis* sp. and *M. breviceaudata*.

NORs confirmed by FISH using the 18S rDNA probe were present in the short arms of pair 6 in all $2n=14$ species and *G. venusta* ($2n=18$). In *M. demerarae* and *M. murina* sites were also detected in the terminal position of the long arms of pair 5 (Fig. 2, a-III e d-III). In *M. emiliae* ($2n=18$) the NOR was positioned on the short arms of pair 7 (Fig. 3d-III), and in *M. touan* in the X and Y chromosomes, although no 18S site was detected in Y (Fig. 3c-III). Only *Monodelphis breviceaudata* exhibited multiple NORs (Fig. 3e-III), whose sites were in the terminal region of the long arms of pair 7 and the short arms of X and Y.

In *D. marsupialis*, both the 18S rDNA probe and silver were detected in three chromosome pairs. In two pairs, the sites were located in the terminal region of the long arms, while in one pair they were bitelomeric (Fig. 3g-III). However, in regards to activity, there was a variation of four to eight markings.

Discussion

In the last decade, advances in systematic and taxonomic studies of the family Didelphidae introduced changes in the taxonomy and nomenclature of several of its taxa (Jansa and Voss 2000, Voss and Jansa 2003, Voss and Jansa 2009, Rossi et al. 2010, Gutiérrez et al. 2010). We used the phylogenetic tree of Jansa and Voss (2014) to map the cytogenetic data of the 18 species we have analysed in order to gain an understanding of chromosome evolution in the group. This work represents the most updated phylogeny of the intergeneric relationships of didelphid marsupials, making our interpretation of the cytogenetic data more integrative than a mere consideration of chromosomal data, and more accurate in light of an independently generated phylogenetic hypothesis.

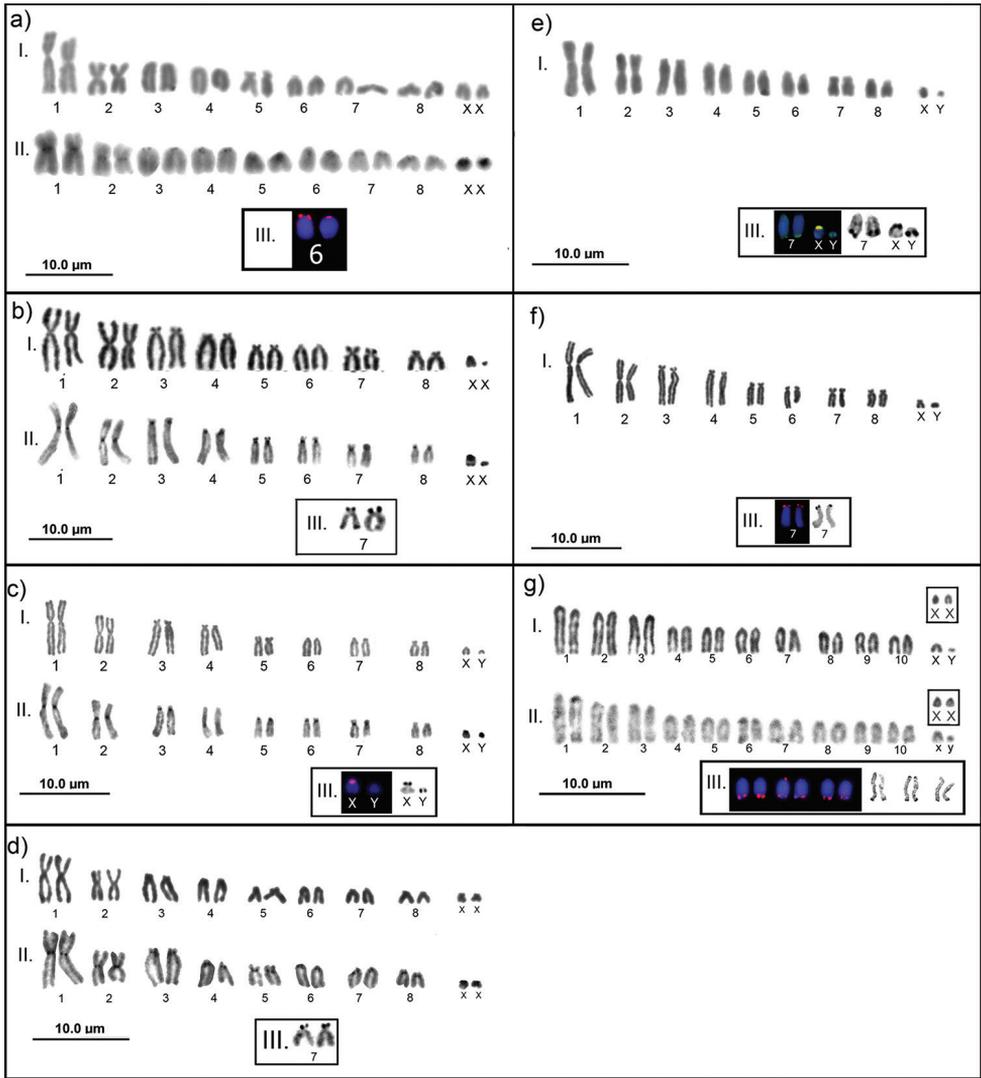


Figure 3. Karyotypes under conventional staining (I), C-band (II), 18S rDNA and Ag-NOR (III), sex chromosomes in the boxes: **a** *Glironia venusta* **b** *Monodelphis* aff. *adusta* **c** *Monodelphis touan* **d** *Monodelphis emiliae* **e** *Monodelphis breviceaudata* **f** *Monodelphis* sp. **g** *Didelphis marsupialis*.

The autosome structure observed here corroborates karyotypic conservation in the diploid number and chromosomal formula (NFa) as previously described in the didelphid species *Didelphis marsupialis*, *Marmosa demerarae*, *Metachirus nudicaudatus*, *Monodelphis touan* (previously named *M. breviceaudata*), *Monodelphis* aff. *adusta* (previously named as *M. cf. emiliae*) and for species of *Marmosops* (Reig et al. 1977, Yonenaga-Yassuda et al. 1982, Casartelli et al. 1986, Hayman 1990, Souza et al. 1990, Palma and Yates 1996, Svartman and Vianna-Morgante 1998, 1999, Carvalho et al. 2002).

Although didelphids are generally considered to have a conserved karyotype, by comparing the karyotypes among different genera, it was possible to associate them with certain species due to the presence of diagnostic characters. For example, *M. demerarae* and *M. murina* differ in their FNa, morphology, and sex chromosome size. In species of the genus *Monodelphis*, morphological variation in chromosomes was restricted to pair 6, which grants an FNa varying between 30 (as observed in *M. aff. adusta*, *M. touan* and, *M. brevicaudata*) and 32 arms (*Monodelphis* sp.). However, the same does not occur for the genus *Marmosops*, in which the five species analysed, present a very similar chromosome macrostructure.

The genus *Marmosa* has a complex taxonomy and recently underwent great taxonomic changes, with all species of *Micoureus*, formerly treated as a separate genus, now considered as a subgenus of *Marmosa*. Considering the taxonomic instability in Didelphidae, with individuals being reclassified, and some complex of species being divided into two or more valid taxa, even purportedly karyotyped species may in fact have their karyotypes still unknown. Thus, our knowledge as to how many and which species among didelphids were karyotyped remains unstable. A revision of the literature for species with reported karyotypes is required.

X chromosome variations

Souza et al. (2013) observed different forms of the X chromosome in *Caluromys phillander*, and our data contribute to show their wide geographic distributions. The acrocentric X are found in northeastern and southeastern Brazil (Fig. 4, localities 16 and 17), as well as in central (Fig. 4, locality 6) and eastern Amazon (Fig. 4, localities 10, 11 and 12). While submetacentric form is located in Venezuela (Fig. 4, locality 14) and areas in the western, central and eastern Amazon (Fig. 4, localities 4, 6, 12 and 15) (Reig et al. 1977, Svartman and Vianna-Morgante 1999, Pereira et al. 2008). Interestingly, both homozygote and heterozygote females were recorded in central Amazonia near Manaus (Fig. 4, locality 6). It is not clear how often this condition is found in natural populations. Indeed, so far, the few heterozygous records for X might be related to the low number of captured and cytogenetically analyzed individuals.

Apparently, there is a likely geographical structure in the distribution of the morphological forms of the X chromosome in *Marmosa murina*, with the metacentric X so far found in the northern and western parts of its distribution, the submetacentric X prevailing in the Amazon basin of Brazil and the acrocentric forms prevailing in the other known localities in central and eastern Brazil (Fig. 5). According to Brito et al (2015), this species is currently under revision and is likely to be split into three species. It remains to be seen if there will be a correspondence between those species and the karyotypic forms depicted here.

Among the Amazonian marsupials analyzed here, the variation in centromere position and heterochromatin patterns of the X chromosome is noteworthy. Souza et al. (2013) suggested that pericentric inversions in the X chromosome of *Caluromys*

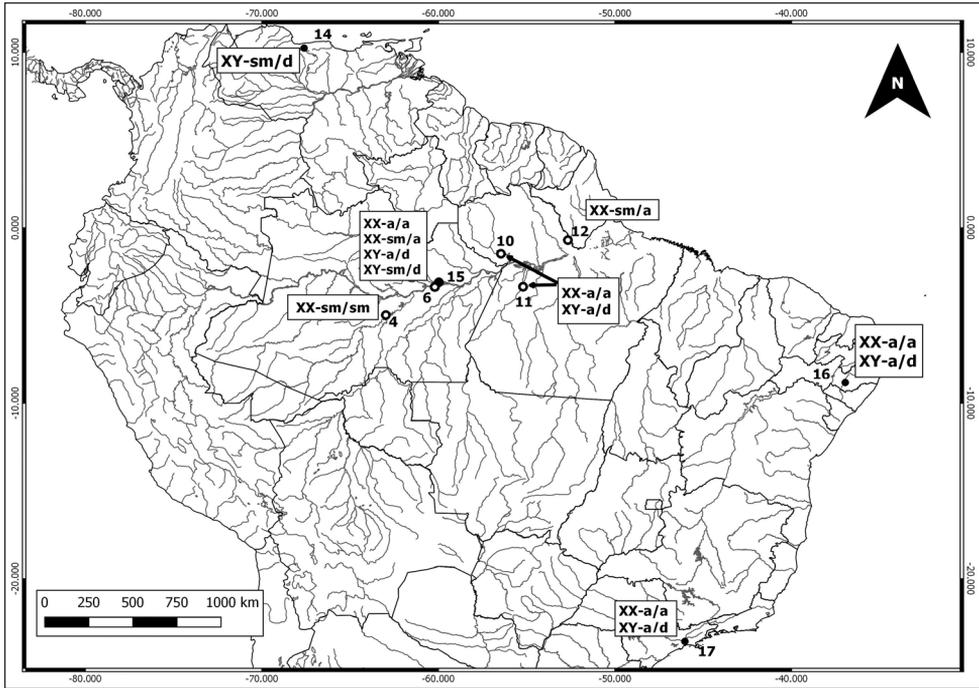


Figure 4. Geographic locations of *Caluromys philander* individuals and its sexual chromosomes morphology data in South America. Literature data represented by empty circles and present work represented by full circles: (● 14) Venezuela, Reig et al. 1977; (○ 4) Purus River; (○ 6) Manaus city, urban área: Federal University of Amazonas’s campus (UFAM); (● 15) Manaus REMAN (Isaac Saba Oil Refinery), present work and Souza et al. 2013; (○ 10) Trombetas River; (○ 11) Tapajós River; (○ 12) Jari River, Souza et al. 2013; (● 16) Pernambuco State, Souza et al. 1990; (● 17) São Paulo State, Svartman and Vianna-Morgante 1999 and Pereira et al. 2008. m=metacentric; sm=submetacentric; a=acrocentric; d=dot-like.

philander altered its morphology, and our results support their findings. In contrast, in *M. murina*, the different morphologies (m, sm, and a) of chromosome X might be due to centromeric shift without the presence of rearrangements. Such reorganization was already observed in other mammals and might be related to three main regions of the chromosome: subtelomeric, proximal and in the boundary between heterochromatin and euchromatin (Rocchi et al. 2012, Burrack and Berman 2012).

Heterochromatin distribution

We observed chromosomal conservatism in the heterochromatin pattern in eight didelphid species: (*C. lanatus*, *G. venusta*, *D. marsupialis*, *M. touan*, *M. aff. adusta*, *M. emiliae*, *G. emiliae* and *M. nudicaudatus*). *C. philander* presented heterochromatic pattern different from the heterochromatic distribution reported in the literature for this

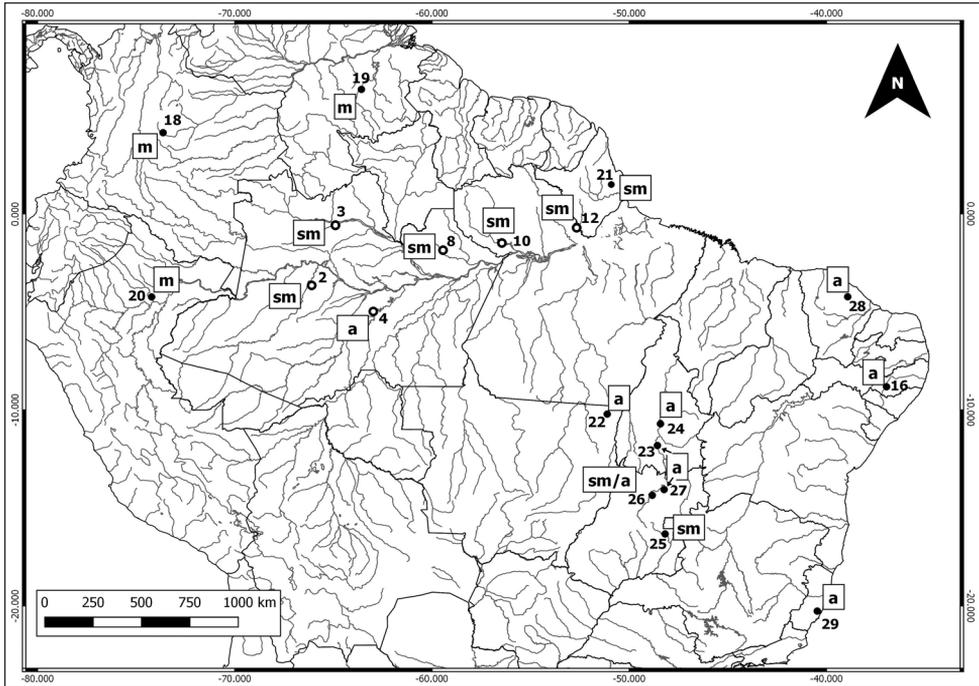


Figure 5. Geographic locations of *Marmosa murina* individuals and its X chromosome morphology data in South America. Literature data represented by empty circles and present work represented by full circles: (● 18) Villa Vivêncio, Colômbia, Hayman and Martin 1974; (● 20) Loreto-Peru, Reig et al. 1977; (● 19) Bolívar, Venezuela, Reig et al. 1977; (○ 3) Negro River; (○ 2) Juruá River; (○ 4) Purus River; (○ 8) Uatumá River; (○ 10) Trombetas River; (○ 12) Jari River; (● 16) Pernambuco State, Souza et al. 1990; (● 21) Tartarugalzinho, Amapá State, Carvalho et al. 2002; (● 22) Vila Rica Mato Grosso State, Pagnozzi et al. 2002; (● 23) UHE Peixe Angical, TO, Pereira et al. 2008; (● 24) Tocantins State, Lima 2004; (● 25) Uruaçu, Goiás State, Carvalho et al. 2002; (● 26) Colinas do Sul, Goiás state; (● 27) UHE Corumbá IV Luziania, Goiás state Pereira et al. 2008; (● 28) Pacoti, Ceará State, Pagnozzi et al. 2002; (● 29) Espírito Santo State, Paresque et al. 2004. m=metacentric; sm=submetacentric; a=acrocentric.

species (Souza et al. 1990, Souza et al. 2013). In *Marmosops* spp., the C-band patterns of the X chromosome are widespread throughout the Amazon basin, but are found in sympatry in the area between the confluences of the Negro-Purus and the Trombetas-Tapajós Rivers, forming pattern 1 to the west and pattern 2 to the east (Table 2). It remains to be seen if there is a correspondence between these patterns with possible cryptic species to be uncovered by broader molecular systematics and morphological studies of these taxa.

Thus, heterochromatin distribution patterns can serve as a cytotaxonomic character, as well as shedding light on chromosomal evolution and regulation of gene expression. However, our results demonstrate that, except for *Marmosops* spp., the other species under study presented little heterochromatin intraspecific variation, including the X chromosome. Thus, this character alone does not allow for distinguishing among

Table 2. Comparative cytogenetic data of the didelphid species analyzed in the present study and those from the literature. In Locality, numbers indicate sampling sites as in the maps of Figures 1, 4 and 5. Karyotypic data: 2n=diploid number; FNa=autosomal arm number; NOR=Nucleolar Organizer Region; p=short arm; q=long arm. Letters identify the X chromosome morphology: m=metacentric; sm=submetacentric; a=acrocentric; d=dot-like. X chromosome C-Band patterns are identified by A= Centromeric heterochromatin; B= Totally heterochromatic except for a terminal euchromatic band; C= Totally heterochromatic except for an interstitial euchromatic band; D= short arm and centromere totally heterochromatic.

| Species | Locality number | 2n | FNa | NORs Pair/arm | 18S rDNA | X/Y | X chromosome C-band | Source |
|--------------------------------|--|----|-----|---------------|----------------|-----------|---------------------|--|
| <i>Caluromys philander</i> | 10; 11; 15; 16 | 14 | 22 | 6p | 6p | a/d | B | Souza et al. 1990; Souza et al. 2013; Present work |
| | 4; 6; 14; 15; 17 | 14 | 22 | 6p | 6p | sm/d | B | São Paulo State, Svartman and Vianna–Morgante 1999, Pereira et al. 2008, Souza et al. 2013, Present work |
| | 12 | 14 | 22 | 6p | 6p | sm/a | B | Souza et al. 2013 |
| <i>Caluromys lanatus</i> | 1 | 14 | 22 | 6p | 6p | sm/– | A. | Present work |
| <i>Marmosa murina</i> | 2; 3; 8; 10; 12; 18; 19; 20; 25; 26 | 14 | 22 | 5q;6p | 5q;6p | (m) sm/ d | A | Hayman and Martin 1974, Reig et al. 1977, Pereira et al. 2008, Carvalho et al. 2002, Present work |
| | 16; 22; 25; 24; 26; 27; 28; 29 | 14 | 22 | 5q;6p | 5q;6p | a/ d | A | Carvalho et al. 2002, Pagnozzi et al. 2002, Lima 2004, Paresque et al. 2004, Pereira et al. 2008 |
| | 4 | 14 | 22 | 5q;6p | 1p; 3p; 5q; 6p | a/d | A | Present work |
| <i>Marmosa demeranae</i> | 2; 3; 4; 5; 6; 7; 9; 10; 11; 12 25, 26 | 14 | 20 | 5q; 6p | 5q;6p | a/a | C | Carvalho et al. 2002, Present work |
| | La Paz, Bolívia | 14 | 20 | – | – | a/a | – | Palma and Yates 1996 |
| | 16 | 14 | 24 | 5q; 6p | – | a/a | – | Souza et al. 1990; |
| | – | 14 | 24 | 5pq; 6p | 5pq; 6p | a/a | – | Svartman and Vianna Morgante 2003 |
| | Rio Grande do Sul | 14 | 24 | 5pq; 6p | – | a/a | – | Carvalho et al. 2002 |
| <i>Marmosops bishopi</i> | 4; 7; | 14 | 24 | 6p | – | m/a | C; D | Present work |
| | 3 | 14 | 24 | 6p | – | m/a | C | Present work |
| <i>Marmosops pinheiroi</i> | 11 | 14 | 24 | 6p | 6p | m/a | C | Present work |
| <i>Marmosops parvidens</i> | 5; 10; 12 | 14 | 24 | 6p | – | m/a | C; D | Present work |
| | 9 | 14 | 24 | 6p | – | m/a | D | Present work |
| <i>Marmosops impavidus</i> | 2 | 14 | 24 | 6p | – | m/a | C | Present work |
| <i>Marmosops pakaraimae</i> | 1 | 14 | 24 | 6p | – | m/a | C | Present work |
| <i>Gracilinanus emiliae</i> | 11; 25; 26 | 14 | 22 | 6p | 6p | m/a | A | Carvalho et al. 2002, Present work |
| <i>Metachirus nudicaudatus</i> | 2; 5 10; 11; 12 | 14 | 20 | 6p | 6p | a/a | A | Present work |

| Species | Locality number | 2n | FN _a | NORs Pair/arm | 18S rDNA | X/Y | X chromosome C-band | Source |
|---------------------------------|-------------------------|----|-----------------|---------------|------------|------|---------------------|--------------------------------------|
| <i>Glironia venusta</i> | 13 | 18 | 20 | 6p | 6p | a/- | A | Fantin e da Silva 2011, Present work |
| <i>Monodelphis touan</i> | 12 | 18 | 28 | Xp | Xp | a/a | A | Present work |
| <i>Monodelphis sp.</i> | 4 | 18 | 32 | 7p | 7p | sm/a | - | Present work |
| <i>Monodelphis aff. adusta</i> | 7 | 18 | 30 | 7p | | a/d | A | Present work |
| <i>Monodelphis emiliae</i> | 7, Juruá River, Acre | 18 | 30 | 7p | | sm/- | A | Patton et al. 2000, Present work |
| <i>Monodelphis brevicaudata</i> | 3; | 18 | 30 | 7q, Xp, Yq | 7q, Xp, Yq | sm/a | - | Present work |
| | Roraima and Pará states | 18 | 30 | Xp | | a/d | | Carvalho et al. 2002 |
| <i>Didelphis marsupialis</i> | 5; 6; 9; 10; 11 | 22 | 20 | 5q;7p;q;8q | 5q;7p-q;8q | a/a | A | Present work |

didelphid populations, although heterochromatin distribution may be an effective character for distinguishing between certain species pairs. This is the case for *M. demerarae* and *M. murina*, with the former presenting larger centromeric heterochromatic blocks than the latter, and between *C. philander* and *C. lanatus*, both with $2n=14$ and $NF=24$, but with distinct heterochromatic patterns.

Nucleolus organizer regions (NORs) and their evolution

The NOR in Didelphidae can be simple or multiple. According to Hsu et al. (1975), the single NOR would be an ancestral character in mammals, with subsequent rearrangements leading to multiple NORs in derived groups. The presence of NOR in sex chromosomes also could be considered a derived character since originally it was present in autosomes and ended up in the X chromosome due to rearrangements such as translocation or transposition. The NOR in *Glironia*, *Monodelphis*, *Caluromys*, *Gracilinanus*, and *Marmosops* is simple. Thus, these genera have a plesiomorphic condition for this character. Conversely, the species of *Didelphis*, *Marmosa* and *Philander* have the derived condition of multiple NORs (Yonenaga-Yassuda et al. 1982, Svartman and Vianna-Morgante 2003).

According to the literature, in *Monodelphis* there are NOR sites on pair 7 and on the X chromosome of *Monodelphis aff. adusta* and *Monodelphis sp.* (Svartman and Vianna-Morgante 1999, Merry et al. 1983, Carvalho et al. 2002). In *M. touan* and *M. brevicaudata* there are simple NORs on the X and Y chromosomes, a condition previously identified in *Monodelphis domestica* Wagner, 1842 (Merry et al. 1983, Pathak et al. 1993). Hsu et al. (1975) reported ribosomal genes in mammal sex chromosomes of the bat species *Carollia castanea*. These authors emphasize that NOR in the X chromosome can generate problems with dosage compensation in mammals.

In the Y chromosome of *M. touan*, FISH did not confirm the marking. This situation was verified in other organisms, where precipitation in the heterochromatic regions took place but could lead to an erroneous interpretation of the distribution of this marker (Schneider et al. 2012). Thus, the marking observed (Fig. 3c III) was not a ribosomal site but a heterochromatic block with silver affinity.

When mapping the NOR character on the phylogenetic tree of Jansa and Voss (2014, fig. 01) (not shown here), we verified that multiple NORs are distributed in two distinct lineages: the first in species of the genus *Marmosa* and the second in species with 22 chromosomes of the genera *Didelphis* and *Philander* Brisson, 1762. The mapping of the simple condition onto the phylogenetic tree depicts a wide distribution for this character, present at the base of the tree (*Caluromys philander*, *C. lanatus*, *Glironia venusta*) and in at least one or more species of the remaining major clades (*Gracilinanus emiliae*, *Marmosops* spp., *Metachirus nudicaudatus*, *Monodelphis touan*, *Monodelphis kunsii*, and *Monodelphis dimidiata*) (Souza et al. 1990, Palma and Yates 1996, Carvalho et al. 2002, Svartman and Vianna-Morgante 2003, Pereira et al. 2008, Souza et al. 2013). This distribution of NOR character on the didelphid phylogeny is thus congruent with the hypothesis advanced by Hsu et al. (1975) that the single NOR is an ancestral state.

When mapping the NOR character on the phylogenetic tree of Pavan et al. (2014) for the genus *Monodelphis*, we verified that *M. emiliae*, *Monodelphis* sp. and *Monodelphis* aff. *adusta* seem to have retained the plesiomorphic condition of a simple NOR. Conversely, this condition became variable in *M. domestica* and in the *M. brevicaudata* species complex, which in addition to the NOR identified in the autosomal pair 7, also presents NORs in both chromosomes of the sex pair, indicating a duplication of this site.

In *M. murina*, intraspecific geographic variation in NORs were detected. Specimens from the Purus River have multiple NORs, those collected in the state of Goiás have simple NOR in the short arms of pair 6 (Palma and Yates 1996, Carvalho et al. 2002) and those from the state of Pernambuco present additional markings in the long arms of pairs 3 and 5 (Souza et al. 1990). Furthermore, both specimens from the Purus River differed from the others regarding sex chromosomes.

Our results indicate geographic variation in NORs for *M. demerarae*. Amazonian specimens analysed did not present ribosomal cistrons in the short arms of the fifth pair, as recorded for specimens from the Atlantic forest in the Rio Grande do Sul and São Paulo states of southern Brazil (Carvalho et al 2002, Svartman and Vianna-Morgante 2003, Svartman 2008). Several studies have shown that considerable genetic variation exists among referred populations of this taxon (Voss and Jansa 2003, Dias et al. 2010, Gutiérrez et al. 2010). Therefore, several nominal taxa previously considered synonyms are now treated as valid species. Currently, *M. demerarae* is considered to occur in south to northern and central Brazil, and to southern Bahia (Gardner 2008, Dias et al. 2010) and *Marmosa paraguayana* Tate 1931 occurs from northern border of Espírito Santo state, south to Rio Grande do Sul, and east to Misiones (Argentina), and eastern Paraguay (Gardner 2008). However, some authors consider it to go as far north as Pernambuco state in northeastern Brazil (Voss and Jansa 2003). Thus, considering the geographic dis-

tribution of this taxon, the 18S rDNA data presented for locations in northern and eastern Brazil possibly belong to specimens of *M. paraguayana*. As such, this character would have a cytotaxonomic value, and rearrangements involving the ribosomal sites could be related to speciation events related to this sister-species pair (Gutiérrez et al. 2010).

In *Didelphis marsupialis* from several Amazonian sites, only NOR activity varied, as was already reported in specimens from the Atlantic forest (Yonenaga-Yassuda et al. 1982, Svartman and Vianna-Morgante 2003).

Conclusion

Dutrillaux (1979) suggested that a small sample size would be inadequate for the knowledge of species karyotypes. Heeding this admonition, we used a relatively large number of individuals for each species analysed to uncover a range of variations that most likely would not have been detected had we used fewer individuals per species. The use of integrative analyses and new methodologies, such as taxonomy, phylogeny, and molecular cytogenetics could improve our understanding of the significance of these chromosomal variations. However, for the Amazon region, a significant limitation for cytogenetic studies is still the restricted collection effort, the vast geographical extent of the region and the difficulty of access to remote areas.

The cytogenetic data presented here shows that didelphid marsupial karyotypes present intraspecific variation in the morphology of sex chromosomes and in chromosomal markers (C-band and NOR) and present some geographic variation in the distribution of these features for several species. Furthermore, there are many areas in the Amazon, including the transition zone between the Amazon and the Cerrado biomes, which do not have cytogenetic records for any didelphid species. This situation seriously undermines our understanding of the significance of the recorded variation, whether it is part of a continuous gradient, or whether it represents intraspecific gradations, or whether it is related to new lineages or cryptic species still uncovered. Thereby, despite the chromosomal stability related to diploid numbers and chromosomal formula in marsupials across continents, didelphids present some intra- and interspecific chromosomal variations, probably related to frequent chromosomal rearrangements. Additional systematic sampling and analyses will be required for a better understanding of the karyotypic evolution of this group.

Acknowledgments

This work was supported by Fundação de Amparo a Pesquisas do Estado do Amazonas (FAPEAM), Coordenação de Aperfeiçoamento Pessoal (CAPES), the SISBIOTA/Rede BIOPHAM (CNPq); and Pró-Amazônia (CAPES). We thank Dra. MC Gross e Dr. CH Schneider who cooperated with research development and G.H. Shepard Jr. for reviewing the English of the manuscript.

References

- Brandão MV, Rossi RV, Semedo TBF, Pavan SE (2015) Diagnose e distribuição geográfica dos marsupiais da Amazônia brasileira. In: Mendes-Oliveira AC, Miranda CL (Eds) Pequenos mamíferos não-voadores da Amazônia brasileira. Sociedade Brasileira de Mastozoologia. Rio de Janeiro, RJ, Brasil, 95–148.
- Brito D, Astua de Moraes D, Lew D, Soriano P, Emmons L (2015) *Marmosa murina*. The IUCN Red List of Threatened Species 2015: e.T40505A22174039. <http://dx.doi.org/10.2305/IUCN.UK.2015-4.RLTS.T40505A22174039.en> [accessed 10 May, 2017]
- Burrack LS, Berman J (2012) Flexibility of centromere and kinetochore structures. *Trends in Genetics* 28(5): 204–212. <https://doi.org/10.1016/j.tig.2012.02.003>
- Carvalho BA, Oliveira LF, Nunes AP, Mattevi MS (2002) Karyotypes of nineteen marsupial species from Brazil. *Journal of Mammalogy* 83(1): 58–70. [https://doi.org/10.1644/1545-1542\(2002\)083<0058:KONMSF>2.0.CO;2](https://doi.org/10.1644/1545-1542(2002)083<0058:KONMSF>2.0.CO;2)
- Casartelli C, Rogatto SR, Ferrari I (1986) Cytogenetic analyses of some Brazilian marsupials (Didelphidae: Marsupialia). *Canadian Journal of Genetics and Cytology* 28: 21–29. <http://dx.doi.org/10.1139/g86-005>
- Dias IMG, Almeida FC, Amato G, De Salle R, Fonseca CG (2010) Delineating geographic boundaries of the woolly mouse opossums *Micoureus demerarae* and *Micoureus paraguayanus* (Didelphimorphia: Didelphidae). *Conservation Genetics* 11: 1579–1585. <http://dx.doi.org/10.1007/s10592-009-9962-5>
- Dutrillaux B (1979) Chromosomal evolution in primates: tentative phylogeny from *Microcebus murinus* (Prosimian) to man. *Human Genetics* 48: 251–314. <http://dx.doi.org/10.1007/BF00272830>
- Fantin C, da Silva MNF (2011) The karyotype of a rare South American marsupial. The bushy-tailed opossum genus *Glirionia* (Didelphimorphia: Didelphidae). *Mastozoologia Neotropical* 18(1): 125–130. <http://www.scielo.org.ar/pdf/mznt/v18n1/v18n1a11.pdf>
- Ford C, Hamerton J (1956) A colchicine hypotonic citrate squash sequence for mammalian chromosomes. *Stain Technology* 31: 247–251. <http://dx.doi.org/10.3109/10520295609113814>
- Gardner AL (2008) *Mammals of South America, volume 1: marsupials, xenarthrans, shrews, and bats*. University of Chicago Press, 669 pp. <https://doi.org/10.7208/chicago/97802262-82428.001.0001>
- Gross MC, Schneider CH, Valente GT, Martins C, Feldberg E (2010) Variability of 18S rDNA locus among *Symphysodon* fishes: chromosomal rearrangements. *Journal Fish Biology* 76: 1117–1127. <http://dx.doi.org/10.1111/j.1095-8649.2010.02550.x>
- Gutiérrez EE, Jansa AS, Voss RS (2010) Molecular systematics of mouse opossums (Didelphidae: *Marmosa*): assessing species limits using mitochondrial DNA sequences, with comments on phylogenetic relationships and biogeography. *American Museum Novitates* 3692: 1–22. <http://dx.doi.org/10.1206/708.1>
- Hayman DL, Martin PG (1974) *Mammalia I: Monotremata and Marsupialia*. In: *Animal cytogenetics*. Gebrüder Borntraeger, Berlin, Germany, 1–110.
- Hayman DL (1990) Marsupial cytogenetics. *Australian Journal of Zoology* 37: 331–349. <http://dx.doi.org/10.1071/ZO9890331>

- Howell WM, Black DA (1980) Controlled silver staining of nucleolus organizer region with a protective colloidal developer: a 1-step method. *Experientia* 36: 1014–1015. <http://doi.org/10.1007/BF01953855>
- Hsu TC, Spirito SE, Pardue ML (1975) Distribution of 18+28S ribosomal genes in mammalian genomes. *Chromosoma* 53: 25–36. <http://doi.org/10.1007/BF00329388>
- Jansa SA, Voss RS (2000) Phylogenetic studies on didelphid marsupials I. Introduction and preliminary results from nuclear IRBP gene sequences. *Journal of Mammalian Evolution* 7(1): 43–77. <http://doi.org/10.1023/A:1009465716811>
- Jansa SA, Baker FK, Voss RS (2014) The early diversification history of didelphid marsupials: a window into South America's "splendid isolation". *Evolution* 68(3): 684–695. <http://dx.doi.org/10.5061/dryad.f1r72>
- Lima JFS (2004) Cariótipos e Regiões Organizadoras de Nucléolo (RON) de *Marmosa* e *Didelphis* (Didelphidae) do estado do Tocantins, Brasil. *Revista Nordestina de Biologia* 18(2): 87–93. <http://periodicos.ufpb.br/ojs/index.php/revnebio/article/view/2610/4380>
- Merry DE, Pathak S, VandeBerg JL (1983) Differential NOR activities in somatic and germ cells of *Monodelphis domestica* (Marsupialia, Mammalia). *Cytogenetics and Cell Genetics* 35: 244–251. <http://doi.org/10.1159/000131875>
- Nagamachi CY, Feldberg E, Pieczarka J, Pereira AL, Silva CEF, Rosa CC, Souza EMS, Pinto JA, da Costa MJR, Malcher SM, Paixão VS, Silva WO (2015) Citogenética de pequenos mamíferos não-voadores da Amazônia brasileira. In: Mendes-Oliveira AC, Miranda CL (Eds) Pequenos mamíferos não-voadores da Amazônia brasileira. Sociedade Brasileira de Mastozoologia, 275–307.
- Pagnozzi JM, Ditchfield AD, Yonenaga-Yassuda Y (2002) Mapping the distribution of the interstitial telomeric (TTAGGG)_n sequences in eight species of Brazilian marsupials (Didelphidae) by FISH and the correlation with constitutive heterochromatin. Do ITS represent evidence for fusion events in American marsupials? *Cytogenetics and Genome Research* 98: 278–284. <http://doi.org/10.1159/000071049>
- Palma RE, Yates TL (1996) The chromosomes of Bolivian didelphid marsupials. *Occasional Papers of Museum Texas Technology University* 162: 1–20. <http://www.nslr.ttu.edu/publications/opapers/ops/OP162.pdf>
- Paresque R, Souza WP, Mendes SL, Fagundes V (2004) Composição cariotípica da fauna de roedores e marsupiais de duas áreas de Mata Atlântica do Espírito Santo, Brasil. *Boletim do Museu de Biologia Mello Leitão* 17: 5–33. http://inma.gov.br/downloads/boletim/arquivos/17/Boletim_17_Artigo1.pdf
- Pathak S, Rønne M, Brown NM, Furlong CL, VandeBerg JL (1993) A high resolution banding pattern ideogram of *Monodelphis domestica* chromosomes. *Cytogenetics and Cell Genetic* 63: 181–184. <http://dx.doi.org/10.1159/000133529>
- Patton JL (1967) Chromosome studies of certain pocket mice, genus *Perognathus* (Rodentia: Heteromyidae). *Journal of Mammalogy* 48(1): 27–37. <https://doi.org/10.2307/1378167>
- Patton JL, da Silva MNF, Malcolm JR (2000) Mammals of the rio Juruá and the evolutionary and ecological diversification of Amazonia. *Bulletin of the American Museum of Natural History* 244: 1–306. [http://dx.doi.org/10.1206/0003-0090\(2000\)244<0001:MOTRJA>2.0.CO;2](http://dx.doi.org/10.1206/0003-0090(2000)244<0001:MOTRJA>2.0.CO;2)

- Pavan S, Jansa SA, Voss RS (2014) Molecular phylogeny of short-tailed opossums (Didelphidae: *Monodelphis*): Taxonomic implications and tests of evolutionary hypotheses. *Molecular Phylogeny and Evolution* 79: 199–214. <http://dx.doi.org/10.1016/j.ympev.2014.05.029>
- Pereira NP, Ventura V, Silva MCJ, Silva DM, Yonenaga-Yassuda Y, Pellegrino CM (2008) Karyotype characterization and nucleolar organizer regions of marsupial species (Didelphidae) from areas of Cerrado and Atlantic Forest in Brazil. *Genetics and Molecular Biology* 31(4): 887–892. <http://dx.doi.org/10.1590/S1415-47572008005000012>
- Pinkel D, Straume T, Gray JW (1986) Cytogenetic analysis using quantitative, high sensitivity, fluorescence hybridization. *Proceedings of the National Academy of Science* 83: 2934–2938. <https://doi.org/10.1073/pnas.83.9.2934>
- Reig OA, Gardner AL, Bianchi NO, Patton JL (1977) The chromosomes of Didelphidae (Marsupialia) and their evolutionary significance. *Biological Journal of Linnean Society* 9: 191–216. <https://doi.org/10.1111/j.1095-8312.1977.tb00265.x>
- Rocchi M, Archidiacono N, Schemp W, Capozzi O, Stanyon R (2012) Centromere repositioning in mammals. *Heredity* 108: 59–67. <https://doi.org/10.1038/hdy.2011.101>
- Rofe R, Hayman D (1985) G-banding evidence for a conserved complement in the Marsupialia. *Cytogenetics and Cell Genetics* 39: 40–50. <https://doi.org/10.1159/000132101>
- Rossi RV, Bianconi GV, Carmignotto AP, Miranda CL (2010) *Ordem Didelphimorphia. Mamíferos do Brasil: Guia de Identificação*. Technical Books, Rio de Janeiro, 19–74.
- Schneider CH, Gross MC, Terencio ML, Artoni RF, Vicari MR, Martins C, Feldberg E (2012) Chromosomal evolution of Neotropical cichlids: the role of repetitive DNA sequences in the organization and structure of karyotype. *Reviews in Fish Biology and Fisheries* 23(2): 201–214. <https://doi.org/10.1007/s11160-012-9285-3>
- da Silva MNF, Rylands AB, Patton JL (2001) Biogeografia e conservação da mastofauna na Floresta Amazônica Brasileira. In: Capobianco JPR, Veríssimo A, Moreira A, Sawyer D, Santos I, Pinto LP (Eds) *Biodiversidade na Amazônia Brasileira: avaliação e ações prioritárias para a conservação, uso sustentável e repartição de benefícios*. Estação Liberdade: Instituto Socioambiental. São Paulo, 110–131.
- Souza MJ, Maia V, Santos JF (1990) Nucleolar organizer regions, G- and C-band in some Brazilian species of Didelphidae. *Revista Brasileira de Genética* 13: 767–775.
- Souza EMS, Silva CEF, Eler ES, da Silva MNF, Feldberg E (2013) Variations of chromosomal structures in *Caluromys philander* (Didelphimorphia: Didelphidae) from the Amazon region. *Genética* 141(1-3): 89–93. <https://doi.org/10.1007/s10709-013-9708-7>
- Sumner AT (1972) A simple technique for demonstrating centromeric heterochromatin. *Experimental Cell Research* 75: 304–306. [http://dx.doi.org/10.1016/0014-4827\(72\)90558-7](http://dx.doi.org/10.1016/0014-4827(72)90558-7)
- Svartman M, Vianna-Morgante AM (1998) Karyotype evolution of marsupials: from higher to lower diploid numbers. *Cytogenetics and Cell Genetics* 82: 263–266. <https://doi.org/10.1159/000015114>
- Svartman M, Vianna-Morgante AM (1999) Comparative genome analysis in American marsupials: chromosomes banding and in-situ hybridization. *Chromosome Research* 7: 267–275. <https://doi.org/10.1023/A:1009274813921>
- Svartman M, Vianna-Morgante AM (2003) Conservation of nucleolus organizer in American marsupials (Didelphidae). *Genética* 118: 11–16. <https://doi.org/10.1023/A:1022986600017>

- Svartman M (2008) American marsupial chromosomes: Why study them? *Genetics and Molecular Biology* 32(4): 675–687. <https://doi.org/10.1590/S1415-47572009005000084>
- Voss RL, Emmons LH (1996) Mammalian diversity in Neotropical lowland rainforests: a preliminary assessment. *Bulletin of the American Museum of Natural History* 230: 1–115. <http://digitallibrary.amnh.org/bitstream/handle/2246/1671/B230a01.pdf?sequence=1>
- Voss RS, Jansa SA (2003) Phylogenetic studies on didelphid marsupials II. Non molecular data and new IRBP sequences: separate and combined analyses of didelphine relationships with denser taxon sampling. *Bulletin of the American Museum of Natural History* 276: 1–82. <http://hdl.handle.net/2246/444>
- Voss RS, Jansa SA (2009) Phylogenetic relationships and classification of didelphid marsupials, an extant radiation of new world metatherian mammals. *Bulletin of the American Museum of Natural History* 322: 1–177. <http://hdl.handle.net/2246/5975>
- Yonenaga-Yassuda Y, Kasahara S, Souza MJ, L'abbate M (1982) Constitutive heterochromatin, G-bands and nucleolus-organizer regions in four species of Didelphidae (Marsupialia). *Genetica* 58: 71–77. <http://dx.doi.org/10.1007/BF00056006>
- Westerman M, Meredith RW, Springer MS (2010) Cytogenetics meets phylogenetics: A review of karyotype evolution in Diprotodontian marsupials. *Journal of Heredity* 101(6): 690–702. <https://doi.org/10.1093/jhered/esq076>
- Wilson DE, Reeder DA (2011) Class Mammalia Linnaeus, 1758. In: Zhang Z-Q (Ed.) *Animal biodiversity: An outline of higher-level classification and survey of taxonomic richness*. *Zootaxa* 3148: 56–60. <https://doi.org/10.1093/jhered/esq076>

Supplementary material I

Voucher specimens

Authors: Carlos Eduardo Faresin e Silva, Rodrigo Amaral de Andrade, Érica Martinha Silva de Souza, Eduardo Schmidt Eler, Maria Nazareth Ferreira da Silva, Eliana Feldberg
Data type: Microsoft Word Document (.docx)

Explanation note: All analyzed specimens were deposited at Mammals Collection in the Instituto Nacional de Pesquisas da Amazônia (INPA); specimens are indicated by species, sampling sites, genus and collector number, followed by INPA collection number (in parentheses) when available.

Copyright notice: This dataset is made available under the Open Database License (<http://opendatacommons.org/licenses/odbl/1.0/>). The Open Database License (ODbL) is a license agreement intended to allow users to freely share, modify, and use this Dataset while maintaining this same freedom for others, provided that the original source and author(s) are credited.

Link: <https://doi.org/10.3897/CompCytogen.v11i4.13962.suppl1>