

CYTOGENETIC ANALYSIS OF THE NEOTROPICAL SPIDER
Nephilengys cruentata (ARANEOMORPHAE,
TETRAGNATHIDAE): STANDARD STAINING, NORs,
C-BANDS AND BASE-SPECIFIC FLUOROCHROMES

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ABSTRACT

The aim of this work is to characterize *Nephilengys cruentata* in relation to the diploid number, chromosome morphology, type of sex determination chromosome system, chromosomes bearing the Nucleolar Organizer Regions (NORs), C-banding pattern, and AT or GC repetitive sequences. The chromosome preparations were submitted to standard staining (Giemsa), NOR silver impregnation, C-banding technique, and base-specific fluorochrome staining. The analysis of the cells showed $2n = 24$ and $2n = 26$ chromosomes in the embryos, and $2n = 26$ in the ovarian cells, being all the chromosomes acrocentric. The long arm of the pairs 1, 2 and 3 showed an extensive negative heteropycnotic area when the mitotic metaphases were stained with Giemsa. The sexual chromosomes did not show differential characteristics that allowed to distinguish them from the other chromosomes of the complement. Considering the diploid numbers found in *N. cruentata* and the prevalence of X_1X_2 sex determination chromosome system in Tetragnathidae, *N. cruentata* seems to possess $2n = 24 = 22 + X_1X_2$ in the males, and $2n = 26 = 22 + X_1X_1X_2X_2$ in the females. The pairs 1, 2 and 3 showed NORs which are coincident with the negative heteropycnotic patterns. Using the C-banding technique, the pericentromeric region of the chromosomes revealed small quantity or even absence of constitutive heterochromatin, differing of the C-banding pattern described in other species of spiders. In *N. cruentata* the fluorochromes DAPI/DA, DAPI/MM and CMA_3 /DA revealed that the constitutive heterochromatin is rich in AT bases and the NORs possess repetitive sequences of GC bases.

Key words: chromosome, Araneae, heterochromatin, secondary constriction, Chromomycin A_3 .

RESUMO

Análise citogenética da aranha neotropical *Nephilengys cruentata* (Araneomorphae, Tetragnathidae): coloração convencional, RONS, bandas C e fluorocromos base-específicos

O objetivo deste trabalho é caracterizar *Nephilengys cruentata* em relação ao número diplóide, à morfologia cromossômica, ao tipo de sistema cromossômico de determinação sexual, aos cromossomos portadores de Regiões Organizadoras de Nucléolo (RONS), padrão de bandas C e seqüências AT ou GC repetitivas. As preparações cromossômicas foram submetidas à coloração convencional (Giemsa), à impregnação pelo nitrato de prata, técnica de obtenção de bandas C e à coloração com fluorocromos base-específicos. A análise das células mostrou $2n = 24$ e $2n = 26$ cromossomos nos embriões e $2n = 26$ nas células ovarianas, sendo todos cromossomos acrocêntricos. O braço longo dos pares 1, 2 e 3 apresentou extensa região heteropictônica negativa quando as metafases mitóticas foram coradas com Giemsa. Os cromossomos sexuais

não mostraram características diferenciais que permitissem distingui-los dos outros cromossomos do complemento. Considerando os números diplóides encontrados em *N. cruentata* e a predominância do sistema cromossômico de determinação sexual do tipo X_1X_2 em Tetragnathidae, *N. cruentata* parece contar com $2n = 24 = 22 + X_1X_2$ nos machos e com $2n = 26 = 22 + X_1X_1X_2X_2$ nas fêmeas. Os pares 1, 2 e 3 mostraram RONS coincidentes com as regiões heteropicnóticas negativas. Utilizando a técnica de obtenção de bandas C, a região pericentromérica dos cromossomos revelou pequena quantidade ou até mesmo ausência de heterocromatina constitutiva, diferindo do padrão de bandas C descrito em outras espécies de aranhas. Em *N. cruentata*, os fluorocromos DAPI/DA, DAPI/MM e CMA_3 /DA revelaram que a heterocromatina constitutiva é rica em bases AT e as RONS apresentam seqüências repetidas de bases GC.

Palavras-chave: cromossomo, Araneae, heterocromatina, constrição secundária, Cromomicina A_3 .

INTRODUCTION

The Araneae order has about 38,000 taxonomically known species distributed in 109 families (Platnick, 2002). Of these approximately 460 species have been studied from the cytogenetic point of view, being 32 belonging to the Neotropical region (Hackman, 1948; Suzuki, 1954; Bole-Gowda, 1958; Brum-Zorrila & Cazenave, 1974; Benavente & Wettstein, 1980; Brum-Zorrila & Postiglioni, 1980; Silva, 1988; Scioscia, 1997; Sbalqueiro *et al.*, 1998; Chen, 1999).

In Araneomorphae (suborder Opisthothelae), which comprises the highest number of cytogenetically studied species, the diploid number varies from $2n = 7$, in *Ariadna lateralis* Karsch 1881, to $2n = 52$, in *Agelena naevia* Comstock 1912 (Wallace, 1909; Suzuki, 1954). In some families of this last infraorder there is a predominance of a certain diploid number, such as Araneidae ($2n = 24$), Gnaphosidae ($2n = 22$), Lycosidae ($2n = 28$), Oxyopidae ($2n = 21$), Pisauridae ($2n = 28$), Salticidae ($2n = 28$), Tetragnathidae ($2n = 24$), and Theridiidae ($2n = 22$), which can represent a more adaptive condition.

The great majority of the cytogenetically described spiders possess acrocentric chromosomes. The sex determination system has been determined in some species of Araneomorphae. About 77% of the species possess the type $X_1X_2/X_1X_1X_2X_2$ (Hackman, 1948; Suzuki, 1951, 1954; Bole-Gowda, 1958; Mittal, 1966; Datta & Chatterjee, 1988; Gorlova *et al.*, 1997; Scioscia, 1997), 9% show the type X/XX (Hackman, 1948; Bole-Gowda, 1958; Suzuki, 1954; Benavente & Wettstein, 1980; Datta & Chatterjee, 1983; Rowell, 1985; Gorlova *et al.*, 1997) and 9% have the type $X_1X_2X_3/X_1X_1X_2X_2X_3X_3$ (Hackman, 1948; Suzuki, 1954; Sharma *et al.*, 1959; Sokolov, 1960, 1962; Datta & Chatterjee, 1983, 1988; Rowell, 1985). Only 3

species possess the type $X_1X_2X_3X_4/X_1X_1X_2X_2X_3X_3X_4X_4$ (Datta & Chatterjee, 1983), 5 species possess the type $X_1X_2X_3Y/X_1X_1X_2X_2X_3X_3$ (Maddison, 1982) and 2 species show the type $X_1X_2Y/X_1X_1X_2X_2$ (Silva, 1988; Sbalqueiro *et al.*, 1998). Furthermore, the sex determination chromosome system has not been identified in 3% of the species.

The family Tetragnathidae possesses $2n = 24 = 22 + X_1X_2$ in the males, and $2n = 26 = 22 + X_1X_1X_2X_2$ in the females, in 14 of the 20 karyotyped species. In the males of other species of this family, the diploid chromosome number can be $2n = 24 = 20 + X_1X_2X_3X_4$ in *Meta segmentata* Chyzer & Kulczynski 1891, $2n = 25 = 22 + X_1X_2X_3$ in *Leucauge celebesiana* Walckenaer 1842, *Leucauge decorata* Blackwall 1864 and *Leucauge tessellata* Thorell 1887, and $2n = 22 = 20 + X_1X_2$ in *Tetragnatha extensa* Linnaeus 1758 and *Tetragnatha* sp. (*Tetragnatha obtusa* C. L. Koch 1837 or *Tetragnatha dearmata* Thorell 1873). In all these species the chromosomes are acrocentric (Hackman, 1948; Suzuki, 1951, 1954; Bole-Gowda, 1958; Sharma *et al.*, 1959, 1960; Sokolov, 1960, 1962; Datta & Chatterjee, 1983, 1988; Gorlova *et al.*, 1995).

The present study was conducted on *N. cruentata* with the purpose of characterizing the karyotype of this species in relation to the diploid number, chromosome morphology, type of sex determination chromosome system, C-banding pattern, and chromosomes bearing the nucleolus organizer regions (NORs), and comparing the karyotype data obtained with those ones of related species described in the literature.

MATERIAL AND METHODS

The chromosome preparations were obtained from 16 subadult specimens (10 males and 6

females) and 7 embryos of *N. cruentata*, which were collected in natural populations at the UNESP, Rio Claro, SP, Brazil.

The embryos preparations were performed according to the methodology described by Webb *et al.* (1978), with some modifications.

The gonadal preparations of subadult specimens were carried out according to the following technique: dissect out the gonads in physiologic solution for insects, transfer the material to colchicin solution (0.16% in physiologic solution for insect), leaving for 2 hours, add a volume of hypotonic solution (tap water) equal to that one of colchicine solution for 15 minutes, place the material in a Carnoy I fixative solution for 60 minutes, macerate the material on the surface of the slide in acetic acid (60%) and dry the slide in a heating metal plate (35-40°C). The standard staining was accomplished using a 3% Giemsa solution, for 13-15 minutes. The C-banding technique was performed according to the methodology described by Sumner (1972), with some modifications. The NOR silver impregnation was made using the method of Howell & Black (1980), with some modifications. The fluorochrome staining was obtained according to Schweizer (1980) technique. The fluorochromes employed were 4-6 diamin-2 phenylindole (DAPI), Mitramycin (MM), Chromomycin A₃ (CMA₃), and Distamycin (DA).

RESULTS

Karyotype description

Mitotic metaphases of *N. cruentata* submitted to the standard staining, C-banding, NOR silver impregnation and fluorochrome technique showed $2n = 24$ in the male embryos (Fig. 1A) and $2n = 26$ in the female embryos and in the oogonials of the young specimens (Fig. 1B). All chromosomes are acrocentric and decrease gradually in size. The sexual chromosomes do not show differential characteristics that have allowed to distinguish them from the other chromosomes of the complement.

Standard staining

The analysis of embryonic and oogonial metaphases evidenced that the chromosomes of the pairs 1, 2 and 3 have a prominent negative heteropycnotic region which extends from the median until the telomeric region of the long arm (Fig. 2A, B). The length of this region can vary among different metaphases of the same specimen, being independent

of the chromosomal condensation degree. Association between two chromosomes by these negative heteropycnotic regions was noted in a great number of mitotic metaphases, indicating the presence of a special kind of chromatin (Fig. 2C). In several mitotic metaphases, the majority of the chromosomes shows negative heteropycnotic short arms (Fig. 1B). Cells in meiotic division were not found in the male and female gonadal cytologic preparations, probably due to the fact that the gametogenesis occurs in a previous instar to those ones of subadults and adults, which were employed in this study.

C-banding

In *N. cruentata*, the majority of the chromosomes does not show C bands in the centromeric region (Fig. 3). Evident C bands occur in the interstitial portion of the long arm in the pairs 1 and 6, in almost all the extension of the long arm in the pairs 5 and 11, and in the telomeric region of the long arm in the pair 10. In the chromosomes of the pair 1, the C bands are intercalated from the median to the telomeric region of the long arm, and are partially coincident with the negative heteropycnotic region. In the chromosomes of the pair 6, the C band is heteromorphic in location (Fig. 3).

Silver nitrate impregnation

In *N. cruentata*, the NORs are coincident with the secondary constrictions, which appear in the pairs 1 and 2 (Fig. 4A, B). Embryonic and oogonial metaphases showed a minimum of four and a maximum of five chromosomes bearing the NORs (Fig. 4C, D). In less condensed metaphases, the chromosomes of the pairs 1 and 2 show more than one NOR (Fig. 4B). Interphasic nuclei show a minimum of one and a maximum of nine nucleoli.

Fluorochromes

In the cells stained with DAPI/MM and analyzed with DAPI filter, the nucleoli in the interphasic nuclei, and the negative heteropycnotic regions in the pairs 1, 2 and 3, are DAPI negative (Fig. 5A). In the metaphasic cells stained with CMA₃/DA, the negative heteropycnotic regions in the pairs 1, 2 and 3 show strong fluorescence, and, additionally, the terminal region of two other chromosomes evidence CMA₃ labels (Fig. 5B). With the fluorochromes DAPI/DA, DAPI positive fluorescence in chromosomal elements of the pairs 1, 5, and 6 was verified in mitotic metaphases (Fig. 5C).

DISCUSSION

Some karyotypic characteristics of *N. cruentata*, i.e., $2n = 24$ in the males and $2n = 26$ in the females, being all the chromosomes acrocentric, are similar to those ones found in the majority of the Tetragnathidae species (Table 1). The accomplished cytogenetic analysis did not allow to establish the type of sex determination chromosome system in *N. cruentata*. However,

considering the diploid numbers obtained in the embryos ($2n = 24$ and $2n = 26$) and in the young females ($2n = 26$), as well as the literature information about the predominant sex determination chromosome system in the species of Tetragnathidae (Table 1), *N. cruentata* probably possesses the system X_1X_2 in the males and $X_1X_1X_2X_2$ in the females. The meiotic cells analysis of *N. cruentata* males will certainly contribute for the establishment of this system.

TABLE 1
Tetragnathidae species analyzed from the cytogenetic point of view, with their respective diploid number ($2n$), sex determination chromosome system (SDS), chromosome morphology (CM) and zoogeographical region (ZR). A = acrocentric.

Species	$2n$ (?)	SDS (?)	SDS (?)	CM	ZR	Reference
<i>Leucauge blanda</i>	24	X_1X_2	$X_1X_1X_2X_2$	24A	Palaearctic	Suzuki, 1954
<i>Leucauge celebesiana</i>	25	$X_1X_2X_3$	$X_1X_1X_2X_2X_3X_3$	25A	Oriental	Datta & Chatterjee, 1983
<i>Leucauge decorata</i>	24	X_1X_2	$X_1X_1X_2X_2$	24A	Oriental	Bole-Gowda, 1958
<i>L. decorata</i>	25	$X_1X_2X_3$	$X_1X_1X_2X_2X_3X_3$	25A	Oriental	Datta & Chatterjee, 1988
<i>Leucauge tessellata</i>	25	$X_1X_2X_3$	$X_1X_1X_2X_2X_3X_3$	25A	Oriental	Datta & Chatterjee, 1983
<i>Meta reticulata</i>	24	X_1X_2	$X_1X_1X_2X_2$	24A	Palaearctic	Hackman, 1948
<i>Meta segmentata</i>	24	$X_1X_2X_3X_4$	$X_1X_1X_2X_2X_3X_3X_4X_4$	24A	Oriental	Datta & Chatterjee, 1983
<i>Meta segmentata</i> ssp. <i>mengei</i>	24	X_1X_2	$X_1X_1X_2X_2$	–	Palaearctic	Sokolov, 1960, 1962
<i>Meta yunohamensis</i>	24	X_1X_2	$X_1X_1X_2X_2$	24A	Palaearctic	Suzuki, 1954
<i>Nephila clavata</i>	24	X_1X_2	$X_1X_1X_2X_2$	24A	Palaearctic	Suzuki, 1951
<i>Nephila clavata</i>	24	X_1X_2	$X_1X_1X_2X_2$	24A	Oriental	Datta & Chatterjee, 1988
<i>Nephilengys cruentata</i>	24	X_1X_2 (?)	$X_1X_1X_2X_2$ (?)	24A	Neotropical	Present Study
<i>Pachygnatha clercki</i>	24	X_1X_2	$X_1X_1X_2X_2$	24A	Palaearctic	Gorlov <i>et al.</i> , 1995
<i>Pachygnatha listeri</i>	24	X_1X_2	$X_1X_1X_2X_2$	24A	Palaearctic	Gorlov <i>et al.</i> , 1995
<i>Tetragnatha andamensis</i>	24	X_1X_2	$X_1X_1X_2X_2$	–	Oriental	Datta & Chatterjee, 1983
<i>Tetragnatha extensa</i>	22	X_1X_2	$X_1X_1X_2X_2$	22A	Palaearctic	Hackman, 1948
<i>T. extensa</i>	24	X_1X_2	$X_1X_1X_2X_2$	–	Palaearctic	Sokolov, 1960
<i>Tetragnatha gracilis</i>	24	X_1X_2	$X_1X_1X_2X_2$	–	Oriental	Datta & Chatterjee, 1983
<i>Tetragnatha japonicola</i>	24	X_1X_2	$X_1X_1X_2X_2$	24A	Palaearctic	Suzuki, 1954
<i>Tetragnatha mandibulata</i>	24	X_1X_2	$X_1X_1X_2X_2$	24A	Oriental	Mittal, 1966
<i>Tetragnatha</i> sp.	24	X_1X_2	$X_1X_1X_2X_2$	–	Oriental	Sharma <i>et al.</i> , 1960
<i>Tetragnatha</i> sp.	24	X_1X_2	$X_1X_1X_2X_2$	24A	Oriental	Mittal, 1966
<i>Tetragnatha</i> sp. (<i>obtusa</i> or <i>dearmata</i>)	22	X_1X_2	$X_1X_1X_2X_2$	22A	Palaearctic	Hackman, 1948
<i>Tetragnatha squamata</i>	24	X_1X_2	$X_1X_1X_2X_2$	24A	Palaearctic	Suzuki, 1954

In all analyzed specimens of *N. cruentata*, the pairs 1, 2, and 3 always show prominent negative heteropycnotic regions. These regions can be heteromorphic in size, in the same metaphase or among different metaphases, can promote association between two chromosomes, and are strongly impregnated by the silver nitrate, evidencing typical characteristics of secondary constrictions related to NORs.

Information about secondary constrictions in the spider chromosomes are scarce in the literature. There are descriptions concerned to these regions for just six species belonging to the families Heteropodidae (presently Sparassidae) (Rowell, 1985), Araneidae (Datta & Chatterjee, 1988), Gnaphosidae (Gorlova *et al.*, 1997), Salticidae (Scioscia, 1997), and Oxyopidae (Chen, 1999). In these species, the secondary constrictions are located in different autosomic pairs and/or in the sexual chromosomes, and there is not a distribution pattern of these regions.

The C band pattern obtained in *N. cruentata*, in other words, the small quantity or even the absence of constitutive heterochromatin in the centromeric region of the chromosomes, the presence of interstitial C bands in some chromosomes, and chromosomic arms almost totally heterochromatic strongly disagree with the pattern obtained in other species of spiders. In general, the chromosomes of spider species submitted to the C-banding technique have revealed the presence of constitutive heterochromatin in the centromeric region of all chromosomes of the complement and absence of interstitial C bands (Brum-Zorrilla & Cazenave, 1974; Brum-Zorrilla & Postiglioni, 1980; Rowell, 1985; Datta & Chatterjee, 1988; Gorlova *et al.*, 1997). Additionally, some of these species possess telomeric C bands in some chromosomes, such as *Nephila clavata* L. Koch 1878 (Tetragnathidae), which shows additional C bands in the distal region of five chromosomes (Datta & Chatterjee, 1988).

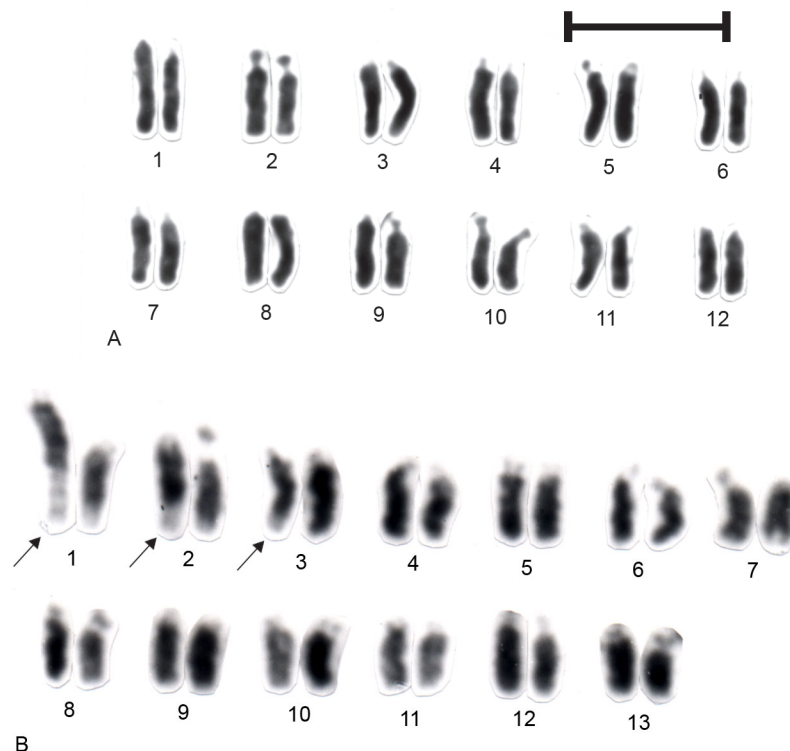


Fig. 1 — Karyotype of *Nephilengys cruentata* conventionally stained. (A) Male embryo, with $2n = 24$. (B) Young female, with $2n = 26$. Arrows = secondary constrictions. Bar: 10 μm .

The small quantity or the absence of constitutive heterochromatin in the centromeric region of the *N. cruentata* chromosomes would be explained by the occurrence of small deletions or by the presence of a special kind of chromatin did not evidence by the C-banding technique employed in this study. The interstitial C bands in the pairs 1 and 6 and the chromosomic arms almost totally heterochromatic in the pairs 5 and 11 would be a consequence of small duplications. In *N. cruentata*, the heteromorphism of interstitial C band location in the chromosomic elements of the pair 6 can be explained as a result of structural rearrangement such as paracentric inversion.

The analysis of *N. cruentata* mitotic metaphases and interphasic nuclei, impregnated by the ion silver shows at least six chromosomes bearing the NORs (pairs 1, 2 and 3) and one to nine nucleoli, respectively. The numerical variability of the NORs and nucleolar material can be a consequence of the differential activity of the NORs. Furthermore, the nucleoli go through association and dissociation periods, during all the cellular cycle, and its number

does not necessarily correspond to the number of chromosomes bearing the NORs (Hsu, 1975; Alberts *et al.*, 2002).

In Araneomorphae, there is little information about the chromosomes bearing the NORs, consequently, there is not a distribution pattern of these regions in the different families. In Dysderidae, one studied species has shown nucleolar material associated to the sexual chromosomes (Benavente & Wettstein, 1980). In Lycosidae, one species has revealed nucleolar material in the terminal region of two bivalents (Wise, 1983). In Tetragnathidae, the distribution pattern of the NORs is being described for the first time in the *N. cruentata* chromosomes.

In *N. cruentata*, the NORs in the pair 1 seem to be intercalated with constitutive heterochromatin, which could have the function of protecting the NORs against mutations and changes, avoiding alterations in the cistrons of rDNA, as proposed by Yunis & Yasmineh (1971), Hsu (1975). NORs adjacent to C bands have also been observed in some insects (Palomeque *et al.*, 1988; Rodríguez-Iñigo *et al.*, 1992; Lorite *et al.*, 1997; Souza *et al.*, 1998).

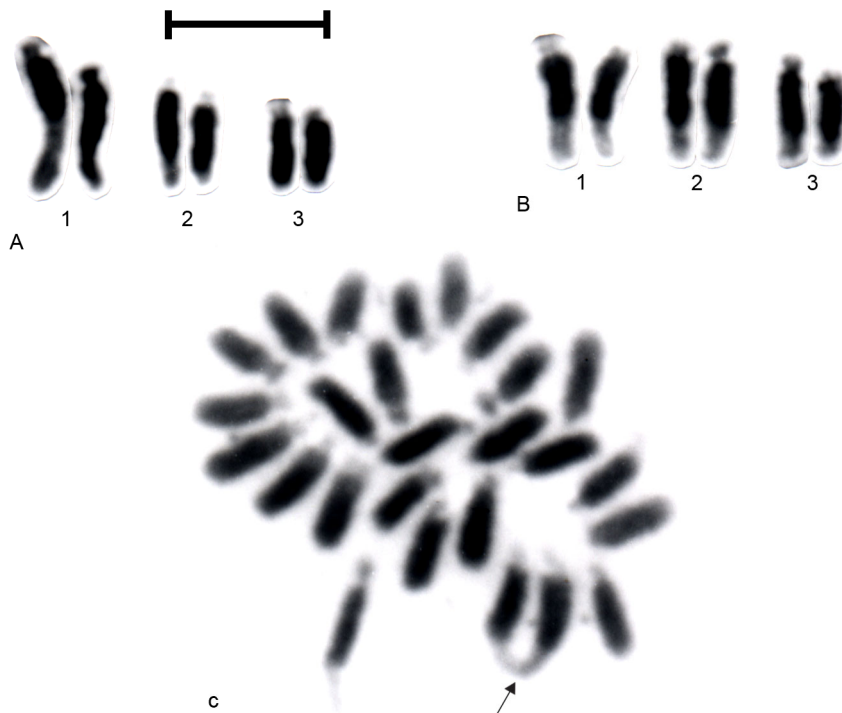


Fig. 2 — Mitotic chromosomes of *Nephilengys cruentata* submitted to the standard staining. (A-B) Pairs 1, 2 and 3 from different metaphases. (C) Metaphase of young female, with $2n = 26$, evidencing association between negative heteropycnotic regions (arrow). Bar: 10 μm .

The mitotic metaphases of *N. cruentata* submitted to the base-specific fluorochromes DAPI/MM and analyzed with DAPI filter have evidenced that the negative heteropycnotic regions in the chromosomal pairs 1, 2 and 3 do not possess highly repeated AT base sequences, since these regions are DAPI negative.

This latter datum was confirmed through the analysis of the results obtained with CMA₃/DA fluorochromes, considering that CMA₃ is GC specific and has positively marked the negative heteropycnotic regions of the pairs 1, 2 and 3. Thus, these results have shown that the NORs in *N. cruentata* possess DNA rich in GC base sequences and poor in AT base sequences. NORs are known to possess GC-rich DNA in several plants (Schweizer, 1976), invertebrates (Schweizer *et al.*, 1983; Lopez-Fernandez *et al.*, 1989; Loreto & Souza, 2000), and vertebrates species (Schmid, 1980).

The presence of CMA₃ positive marks in the terminal region of two other chromosomes of *N.*

cruentata would be related to the inactive NORs, considering that these regions were not evidenced by the silver nitrate impregnation, which marks only active NORs in the preceding interphasis according to Miller *et al.* (1976), Howell (1977) and Schwarzacher *et al.* (1978), or would represent a special kind of chromatin, rich in GC bases, but not evidenced by the C-banding technique employed in the present study.

The positive marks obtained in the *N. cruentata* chromosomes with DAPI/DA fluorochromes are partially coincident with the C-bands. In the pair 1, this heterochromatin rich in AT bases seems to precede the negative heteropycnotic region. In the pair 5, the heterochromatic long arm has revealed a rich AT base sequence and the interstitial heterochromatin of the pair 6 has shown a highly repeated AT base sequence.

Up to this moment, there is no cytogenetic analysis in Araneae using base-specific fluorochromes.

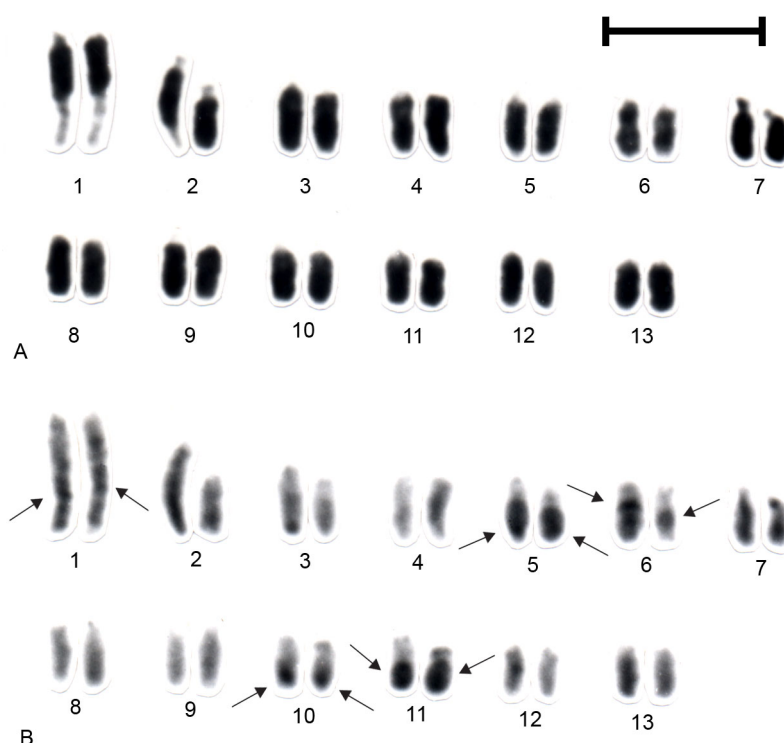


Fig. 3 — Karyotype of *Nephilengys cruentata* female, with $2n = 26$. (A) Standard staining. (B) The same karyotype showed in A submitted to the C-banding technique. Arrows = C bands. Bar: 10 μm .

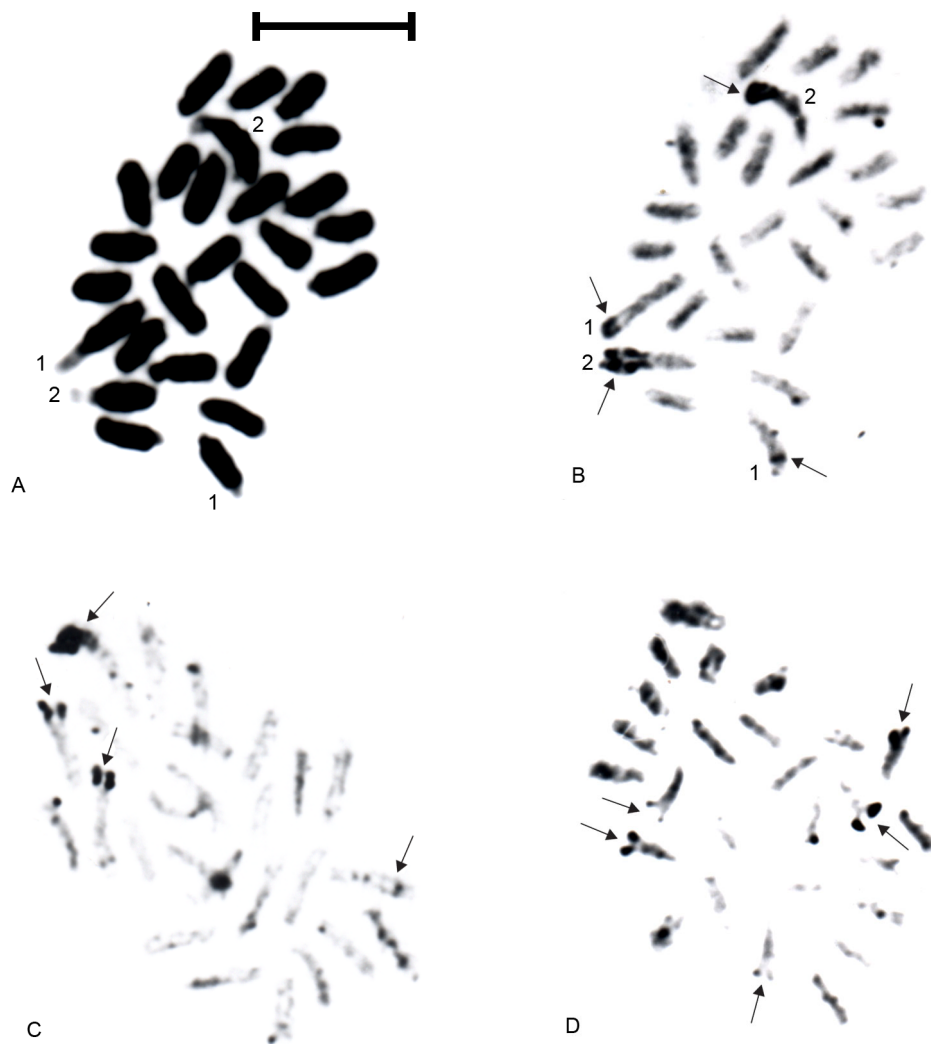


Fig. 4 — Mitotic metaphases of *Nephilengys cruentata* female, with $2n = 26$. (A) Standard staining. (B) The same metaphase showed in A submitted to the silver nitrate impregnation. (C) With 4 NORs. (D) With 5 NORs. Arrows = NORs. Bar: 10 μ m.

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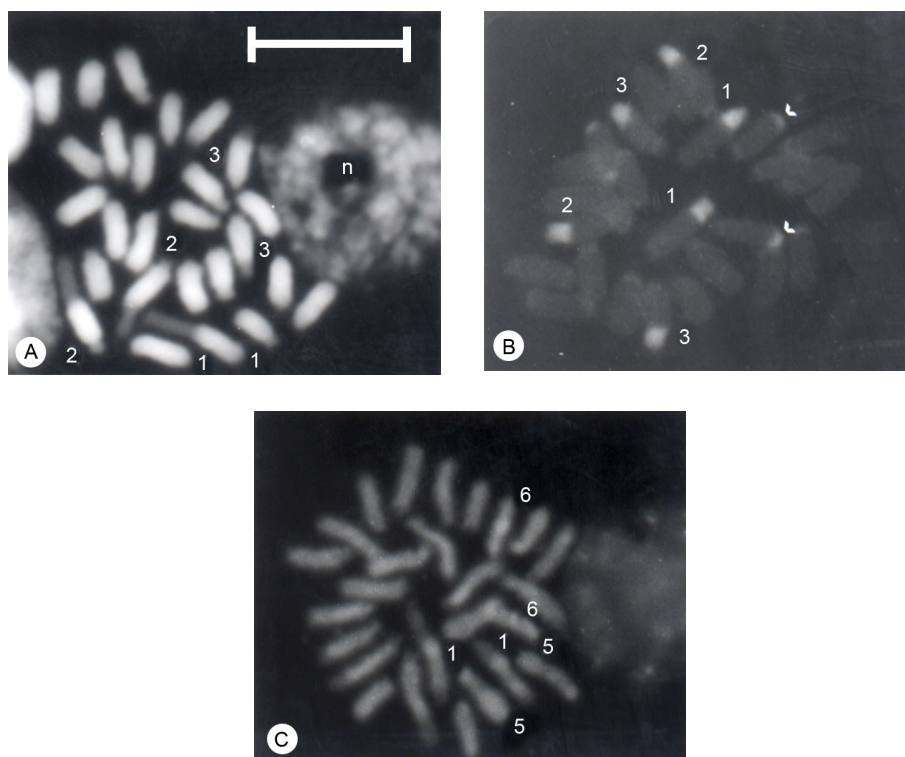


Fig. 5 — Oogonial cells of *Nephilengys cruentata* submitted to the base-specific fluorochromes. (A) Interphasic nuclei and mitotic metaphase with $2n = 26$ stained with DAPI/MM, evidencing nucleoli (n) and DAPI negative chromosomal regions in the pairs 1, 2 and 3. (B) Metaphase with $2n = 26$ submitted to the CMA_3/DA , showing CMA_3 positive regions in the pairs 1, 2 and 3, and in an additional chromosomal pair (arrows). (C) Metaphase with $2n = 26$ stained with DAPI/DA, exhibiting DAPI positive regions in the pairs 1, 5 and 6. Bar: 10 μm .

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