A chromosomal study on a Lebanese spittlebug *Philaenus arslani* (Hemiptera: Auchenorrhyncha: Aphrophoridae)

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Abstract. The meadow spittlebug genus *Philaenus* (Auchenorrhyncha: Aphrophoridae) is known to display marked colour polymorphism. This study presents the results of a karyotype analysis of *P. arslani* from Lebanon using conventional chromosome staining, C-banding, fluorescent banding using base-specific fluorochromes (CMA₃ and DAPI) and AgNOR-staining. This species has 2n = 18 + neo-XY, and differs from *P. spumarius* both in the number of chromosomes and sex chromosome system. During meiosis, the neo-XY bivalent is clearly heteromorphic being the largest in the complement. Furthermore, sex chromosomes show marked differences in C-banding pattern. The NOR-bearing chromosomes are the first and one of the middle-sized pairs of autosomes. NORs are G-C rich. Furthermore, some blocks of constitutive heterochromatin on the sex chromosomes are also G-C rich. All other C-bands are DAPI or DAPI/ CMA₃ positive, thus containing A-T rich DNA. The significant difference in the karyotype of *P. arslani* and *P. spumarius* indicates chromosomal transformations during the evolution of the genus *Philaenus*.

INTRODUCTION

Until quite recently the genus *Philaenus* Stål, 1864 was considered to include three species only: the holarctic *P. spumarius* (Linnaeus, 1758), the Mediterranean *P. signatus* Melichar, 1896 and *P. tesselatus* Melichar, 1889. The latter, of unclear taxonomic status, is regarded also as a synonym of *P. spumarius* (see Drosopoulos & Quartau, 2002). Recent studies increased the number of *Philaenus* species to 8 and all of the newly described species were discovered in the Mediterranean (Drosopoulos & Asche, 1991; Loukas & Drosopoulos, 1992; Abdul-Nour & Lahoud, 1995; Drosopoulos & Remane, 2000; Remane & Drosopoulos, 2001; Drosopoulos & Quartau, 2002). Of these species, *P. spumarius*, *P. tesselatus*, *P. loukasi* Drosopoulos and Asche, 1991 and *P. arslani* comprise the "spumarius" group, whereas *P. tarifa* Remane and Drosopoulos, 2001, *P. signatus*, *P. italosignus* Drosopoulos and Remane, 2000 and *P. maghresignus* Drosopoulos and Remane, 2000 belong to the "signatus" group, however this division is still ambiguous (Loukas & Drosopoulos, 1992; Drosopoulos & Remane, 2000; Drosopoulos, 2003). *P. arslani* was described from Lebanon, where it occurs in mountains at altitudes between 1000–2000 m. This species is associated with arid plants such as *Eryngium*, *Echinops*, *Carduus*, *Cirsium*, *Cistus*, and expresses only one colour morph named *populi* (Abdul-Nour & Lahoud, 1995). Quite recently, populations of this species were also found in SE Turkey and in Iraq (see Drosopoulos, 2003).

Up to now the only *Philaenus* species studied cytogenetically, *P. spumarius*, has 2n = 22 + XX/0 (Boring, 1913; Kurokawa, 1953; Kuznetsova et al., 2003). Some additional information on the karyotype of *P. spumarius* was also provided by several methods of differential chromosome staining (C-, AgNOR- and fluorescent-banding) (Kuznetsova et al., 2003).

The genus *Philaenus* is known to show marked colour polymorphism. The nature and origin of this polymorphism and its possible contribution to the evolution of reproductive isolation and sympatric speciation have been extensively documented for *P. spumarius* (for an overview of the literature, see Halkka & Halkka, 1990; Stewart & Lees, 1996; Drosopoulos, 2003). Alteration of the karyotype may play a substantial role in the process of speciation (King, 1993). Therefore, an interesting question is whether this is the case for the genus *Philaenus*, which is considered to be one of the best insect genera for evolutionary studies on polymorphism and speciation (Drosopoulos & Quartau, 2002).

This paper is part of a project exploring the cytogenetics of the genus *Philaenus* and presents the karyotype of another representative of this group, *P. arslani*, described from Lebanon, which as far as is known does not show colour polymorphism (Abdul-Nour & Lahoud, 1995).

MATERIAL AND METHODS

Specimens of *P. arslani* were collected by H. Abdul-Nour in the mountains of Lebanon, in August 2004, from Lassa (Kesrouane), altitude: 1200 m and in June–July 2005 from Mazzraat Kfar Zeibane (Kesrouane), altitude: 1600 m. Adult specimens were fixed in an ethanol: acetic acid fixative (3:1, 96% ethanol : glacial acetic acid). Testis structure of 18 males and ovary...
structure of 4 females were studied primarily in terms of the number of testicular follicles and ovarioles. The chromosomal study was based on spermatogenesis in 18 males. Chromosome spread preparations of mitotic and meiotic cells obtained from testicular follicles were made as previously described by Kuznetsova et al. (2003). The standard Feulgen-Giemsa procedure (Grozeva & Nokkala, 1996) was used for conventional staining. Silver staining of nucleolar organizing regions (NORs) was performed following the technique of Howell & Black (1980). Some slides were C-banded to visualize constitutive heterochromatin according to Sumner (1972). Chromomycin A₃ (CMA₃) and 4′,6′-diamidino-2-phenylindole (DAPI) staining of heterochromatin indicated by CMA₃ bright/DAPI dark sites (Fig. 14, 15). These CMA₃ positive sites were argen- tophilic demonstrating that these chromosomes bear NORs, containing actively transcribed rDNA genes. Furthermore, silver stained nucleolar remnants were seen attached to these chromosomes during meiosis up to diakinesis (Figs 16, 17).

Mitotic complement

Male karyotype of *P. arslani* consists of 20 holokinetic chromosomes. In mitotic complements, a pair of very large chromosomes, one clearly larger than the other, represent sex chromosomes (Figs 3, 4). The larger sex chromosome is probably an X chromosome and the smaller a Y chromosome (see Discussion).

Course of meiosis

Spermatocyte pachytene cells showed two separate positively heteropycnotic bodies of differing size representing the sex chromosomes (Fig. 5). Diplotene, diakinesis and metaphase I revealed 10 bivalents indicative of 2n = 20 (Figs 6–9). The nine autosomal bivalents gradually decreased in size. Sex chromosome pair was significantly larger than the largest autosomal bivalent and clearly heteromorphic. Each of the bivalents normally formed the only chiasma, however in some cells two or very rarely three chiasmata could be seen in the sex bivalent and in two or even three larger autosomal bivalents. The chiasmata were generally terminally located except for bivalents with three chiasmata, in which one chiasma was interstitial (Figs 6, 7). The anaphase I showed two groups of chromosomes, 9A + X and 9A + Y respectively, segregating to opposite poles (Fig. 10). Thus, the male chromosome complement is suggested to be 2n = 18 + XY.

C- and fluorescent CMA₃/DAPI-banding, AgNOR-staining

C-banding of mitotic chromosomes revealed small C-bands on the telomeres of autosomes. In contrast to autosomes, C-banding showed prominent heterochromatic regions on both telomeres of the putative X chromosomes and on one of telomeres of the putative Y chromosome. On the X chromosome, one of the bands was about twice as large as the other. On the Y chromosome, telomeric heterochromatin consisted of two closely-spaced bands (Figs 3, 4). The banding pattern on the X and the Y chromosomes was always visible also in meiosis, independent of their degree of condensation. Telomeric C-bands of the X chromosome tended to be attracted together, in which case the X very often formed a ring in mitosis (Fig. 3), whereas in meiotic prophase the XY bivalent produced an intricate loop (Figs 6, 7). Staining of C-banded chromosomes by fluorochromes revealed that *P. arslani* heterochromatin contained both A-T and G-C rich DNA (Figs 11–15). In meiosis, telomeric heterochromatin of some of autosomes was brightly coloured after DAPI/CMA₃ staining, evidence that A-T and G-C bases are equally represented at this site (Figs 12, 13). The largest autosomal pair and one of the middle-sized autosomal pairs (6th or 7th) carried only G-C rich heterochromatin indicated by CMA₃ bright/DAPI dark sites (Fig. 14, 15). These CMA₃ positive sites were argentophilic demonstrating that these chromosomes bear NORs, containing actively transcribed rDNA genes. Furthermore, silver stained nucleolar remnants were seen attached to these chromosomes during meiosis up to diakinesis (Figs 16, 17).
The telomeric C-band of the Y chromosome and one terminal band of the X chromosome were brightly fluorescent both after DAPI and CMA3 staining, showing that they are A-T and G-C rich. The other telomeric C-band of the X chromosome displayed a DAPI bright/CMA3 dark fluorescent pattern (Figs 12, 13).

DISCUSSION

On the basis of the morphology of male and female internal reproductive organs, several characters have been identified, including the number of follicles and ovarioles, which may be useful taxonomically as well in understanding the phylogenetic traits of the Auchenorrhyncha (Emelyanov & Kuznetsova, 1983; Kuznetsova, 1985; Kirillova, 1989; Bednarczyk, 1993; Kuznetsova et al., 1998; D’Urso et al., 2005). For the family Aphrophoridae, there is some data on the morphology of testes and ovaries, mainly the number of testicular follicles and ovarioles, for as few as 4 species, including P. spumarius (see Emelyanov & Kuznetsova, 1983; Kuznetsova et al., 2003). These results demonstrate considerable variability in the number of testicular follicles (12–35) and ovarioles (11–20) in the family. Moreover, follicles and ovarioles slightly vary in number between specimens and even between different testes and ovaries of a specimen, as in P. arslani (this study). In this species, between
10 and 12 follicles and from 16 to 19 ovarioles were found in the 18 males and 4 females studied. In *P. spuriosus*, there are 11–13 follicles and 10–12 ovarioles (Ivanov, 1926; Kuznetsova et al., 2003). Hence, congeneric species share a similar testis structure, but differ in the number of ovarioles. However, data for other *Philaenus* species are required before making inferences.

Previous cytogenetic reports on the family Aphrophoridae, including 22 species from 9 genera, have shown diploid chromosome numbers ranging from 16 to 32 (in
female) and the occurrence of two sex chromosome determination systems, XX/X0 and XX/XY (see Halkka, 1959; Kirillova, 1986). The representatives of the family, similar to other auchenorrhynchan groups, have holokinetic chromosomes, a chiasmatic pre-reductional male meiosis with predominantly 1–2 chias mata per bivalent, and fusions/fissions as the basic mode of karyotype evolution, fusions of chromosomes being apparently more common than fissions (Halkka, 1959, 1964; Kuznetsova et al., 2003).

In spite of the high species diversity in the genus Philaenus, only the karyotype of P. spumarius has been studied so far. This species has 2n = 22 + XX/X0 (Boring, 1913; Kurokawa, 1953; Kuznetsova et al., 2003). Furthermore, the karyotype of this species contains two CMA3 positive NORs, associated with the first and one of the middle-sized (6th or 7th) autosome pairs respectively, and very small C-bands located on the telomeres of both autosomes and the X chromosome (Kuznetsova et al., 2003). The karyotype of P. arslani reported here is distinctly different in chromosome number (2n = 18 + neoXY), sex chromosome system (XX/XY), and in the greater amount of sex chromosome heterochromatin.

In contrast to the sex chromosomes, autosomes of both species have only dot-like telomeric C-bands. An important point is that P. arslani and P. spumarius have a similar number and location of NORs, suggesting that the NOR bearing chromosomes were not involved in the rearrangements. The fact that NOR associated heterochromatin stains brightly with CMA3 in both P. spumarius and P. arslani indicates that it is G-C rich. This fluorescence pattern of NOR seems to be a general feature of the hemipteran genome, since it has been described in the great majority of taxa examined so far (Nechaeva et al., 2004; Cattani et al., 2004; Criniti et al., 2005; Lanzone & De Souza, 2006). Noteworthy is that P. spumarius and P. arslani differ significantly in the amount and composition of heterochromatin. In P. spumarius, heterochromatin is only found in NOR bearing chromosomes, whereas P. arslani displays a considerable amount of heterochromatin, which contains both G-C and A-T rich regions. In this species, heterochromatin in the X and Y is both DAPI and CMA3 bright, whereas a telomeric band of the X is DAPI bright/CMA3 dark.

The low number of chiasmata seems to be a standard pattern in holokinetic bivalents (Halkka, 1964). Nokkala et al. (2004) analyzed chiasma formation and distribution in holokinetic bivalents of a psyllid species, Beopalma foersteri (Flor) (Sternorrhyncha: Psylloidea). These authors conclude that more than two chiasmata in a holokinetic bivalent must necessarily obstruct the regular course of meiosis and result in the elimination of cells of this sort. In P. arslani, two and occasionally three chiasmata were observed in separate bivalents. Very rarely, three and even four chiasmata are likewise found in larger bivalents of P. spumarius (Kuznetsova et al., 2003). The fate of cells with multichiasmatic bivalents remains, however, unknown in both species.

Comparison of karyotypes of both Philaenus species indicates that the XY system of P. arslani is formed by neo-sex chromosomes. We suggest that the karyotype of P. arslani has been derived from that of P. spumarius by means of two fusions, the first between two pairs of autosomes and the second between an autosome and the X chromosome. As a consequence, the putative neo-Y chromosome and the autosomally derived part of the putative neo-X chromosome of P. arslani appear similar in size. They are also similar in that each has a C-band on one of the telomeres. Similar to sex chromosomes with standard kinetic structure (monocentric chromosomes), holokinetic sex chromosomes quite often contain a considerable amount of heterochromatin (Papeschi, 1995; Grozeva & Nokkala, 2001; Criniti et al., 2005; Kuznetsova et al., 2007), and this is also true for P. arslani. A large C-heterochromatic segment situated on a telomere of the putative X chromosome consists probably mainly of an original X chromosome of the ancestor. It is noteworthy that a very similar pattern occurs in Neophilaenus lineatus Linnaeus and N. exclamationis Thunberg of the same family. These closely related species have 2n = 28 + XX/X0 and 2n = 18 + XX/XY, respectively. In N. exclamationis, the X and Y chromosomes are very large, the X markedly exceeding the Y in size and carrying a heterochromatin segment. It is speculated that the heterochromatin segment represents a former X chromosome (Halkka, 1959). At present, with data available for only two species, little can be inferred about the chromosomal evolution of the genus Philaenus. However, the significant difference in the karyotype of P. arslani and P. spumarius indicates many chromosomal transformations during the evolution of the genus Philaenus, including structural rearrangements and heterochromatinization. We suggest that data for other Philaenus species will reveal further differentiation at the chromosomal and anatomical levels, giving additional insight into the taxonomy and evolution of this speciose genus.

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