Acta Theriologica 45, Suppl. 1: 59–67, 2000. PL ISSN 0001-7051

Reassessment of the range of the Drnholec race: studies on meiosis in *Sorex araneus* hybrids

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Fedyk S., Banaszek A., Chętnicki W., Cichomska A. and Szałaj K. A. 2000. Reassessment of the range of the Drnholec race: studies on meiosis in *Sorex araneus* hybrids. [In: Evolution in the *Sorex araneus* group: Cytogenetic and molecular aspects. J. B. Searle and J. M. Wójcik, eds]. Acta Theriologica 45, Suppl. 1: 59–67.

The Stobnica race (gm, hi, ko, np) and Drnholec race (gm, hi, ko, nr) are difficult to distinguish and so studies were carried out to determine which is prevalent in eastern Poland. Meiotic studies on hybrids showed that it is the Drnholec race rather than the Stobnica race that makes contact with the Białowieża, Guzowy Młyn and Popielno races.

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Key words: Sorex araneus, hybrid zone, chromosome races, meiosis

Introduction

Cytogeneticists have assumed that centric fusions and whole arm reciprocal translocations are the basis of the phenomenal differentiation of *Sorex araneus* karyotypes. Over 50 chromosomal races have been described up to now (Zima *et al.* 1996). In order to properly establish the chromosomal races of *Sorex araneus*, a method for accurate identification of chromosome arms has been essential. Once differential staining techniques (Q- and G-banding) were applied to common shrew chromosomes (Halkka *et al.* 1974, Fredga and Nawrin 1977), identification of all the chromosome arms became possible. In fact, it is easy to discriminate large and medium size (a-l) chromosome arms (see Searle *et al.* 1991). However, the smallest arms (m-r) can be misidentified.

The chromosome arms p and r are the most difficult to discriminate. Both have one dark band close to the centromere, and in the acrocentric state, they can additionally be mistaken for the Y₁ chromosome (Searle *et al.* 1991). Theoretically, arm r should be smaller than arm p, but this difference in size can only be visible in the best preparations.

Particular precaution is necessary when two very similar metacentrics, that differ in p and r chromosome arms, occupy neighbouring territories. Two contiguous chromosomal races, differing only in nr and np metacentrics, have been described

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in Poland (Fig. 1). One of them, the Stobnica race (gm, hi, ko, np), was described from western Poland (Wójcik 1986) and the shrews with the same specific metacentrics were described as race II from north-eastern Poland (Fedyk 1986, Fedyk and Leniec 1987). The Drnholec race (gm, hi, ko, nr) was described in southern Moravia (Fig. 1), and its range was originally limited to Bohemia and Moravia (Zima and Kral 1985). Zima et al. (1988) suggested that the range of Drnholec race extended northwards, to southern Poland, but Wójcik (1993) considered it limited to the very southernmost part of the country. Wójcik (1993) described the Stobnica race as having a more northern and eastern distribution within Poland. Towards the east, Wójcik (1993) considered that it made contact with the Białowieża, Popielno, Guzowy Młyn and Łęgucki Młyn races, all of which belong to the East European Karyotypic Group (EEKG) as opposed to the Stobnica race which is a member of the West European Karyotypic Group (WEKG) (see Fig. 1). The hybrid zones between the Stobnica and Białowieża races (Fedyk et al. 1993) and the Stobnica and Legucki Młyn races have been described (Fedyk et al. 1991, Szałaj et al. 1995, Fedyk 1995). However, subsequent to Wójcik's distributional

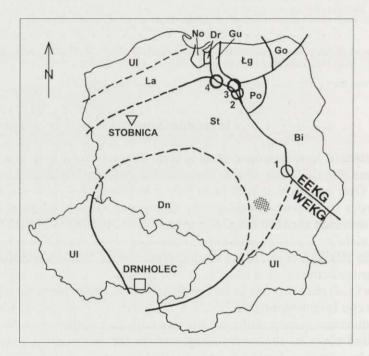


Fig. 1. Distribution of chromosomal races of *Sorex araneus* in Poland (according to Wójcik 1993 and Fedyk 1995), Czech Republic and Slovakia (according to Lukáčová *et al.* 1994). Type localities for the Stobnica and Drnholec races (triangle and square, respectively), and populations studied (circles) are shown. Codes for races: Ul – Ulm, No – Nogat, Dr – Drużno, Gu – Guzowy Młyn, Łg – Łęgucki Młyn, Go – Gołdap, La – Laska, Po – Popielno, Bi – Białowieża, St – Stobnica, Dn – Drnholec. Populations studied: 1 – Plebanka, 2 – Wykno, 3 – Czarny Piec, 4 – Mycyny. Shaded area: isolated population of the Laska race in Świętokrzyskie Mts (Wójcik 1993).

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study, it was shown that shrews of the Drnholec race were found to hybridize with the Białowieża race in south-eastern Poland, hence the range of the Drnholec race was extended to the east (Szałaj *et al.* 1996). Therefore, at least in some parts of contact zone between the races of WEKG and EEKG in Poland, the Drnholec race, not the Stobnica, is the representative of the WEKG races. Given the difficulty in distinguishing nr and np, the occurrence of the Stobnica race in other sections of the contact zone between the WEKG and EEKG races has become doubtful and the aim of the present study is to establish whether it is the Drnholec or Stobnica race that occurs in north-eastern Poland.

Material and methods

We conducted meiotic studies on hybrids collected from four sites in north-eastern Poland from three different hybrid zones between the EEKG and WEKG races. The races involved were known to be the Drnholec (Dn) and Białowieża (Bi) races at Plebanka (site 1 in Fig. 1) (Szałaj *et al.* 1996). In the other sites we were aiming to determine whether the WEKG race was the Stobnica (St) or Dn race. The EEKG races were the Guzowy Młyn (Gu) race at Wykno and Mycyny (sites 2 and 4 in Fig. 1) and the Popielno (Po) race at Czarny Piec (site 3 in Fig. 1). The karyotypes of the F₁ hybrids expected in these hybrid zones are shown in Table 1. It can be seen that the meiosis I configurations would differ between Po × Dn and Po × St F₁ hybrids and between Gu × Dn and Gu × St hybrids.

In an area where there is Robertsonian polymorphism, identification of hybrids with nr or np is more difficult. But the individuals with acrocentrics, instead of one of the metacentrics forming the meiotic complex, can be useful. In such a case a chain-of-five (CH_V) arises instead of a ring-of-four (R_{IV}) configuration when the hybrids possess chromosome nr, whereas in hybrids with the np metacentric, polymorphism of gm or mn divides the CH_{VI} configuration into two complexes (CH_{IV} and CH_{III}), and polymorphism of gr or np chromosomes shortens the CH_{VI} complex to CH_V with an acrocentric bivalent excluded from the chain (Table 1).

Meiotic and mitotic preparations were made according to standard techniques (Verma and Babu 1989). At least 20 good diakineses were analysed from each of four hybrids studied.

Gametes of WEKG races Gametes of EEKG races	Drnholec (Dn) hi, ko, gm, nr, p, q	Stobnica (St) hi, ko, gm, np, r, q
Białowieża (Bi)	o/ok/ki/ih/hn/nr/ rg/gm/mp /p,	o/ok/ki/ih/hn/np/ pm/mg/gr /r
hn, ki, gr, mp, o, q	qq	qq
Popielno (Po)	o/ok/ki/ih/hq/q,	o/ok/ki/ih/hq/q,
hq, ki, gr, mn, o, p	rn/nm/mg/gr,	r/rg/gm/mn/np/ p
	рр	
Guzowy Młyn (Gu)	hi hi, ko ko,	hi hi, ko ko,
hi, ko, gr, mn, p, q	rn/nm/mg/gr,	r/rg/gm/mn/np/p,
	pp , qq	qq

Table 1. Karyotypes expected in F_1 hybrids between the EEKG races and the Drnholec or Stobnica races (WEKG) in eastern Poland. Boldface indicates the position of chromosomes in meiotic complexes in relation to the presence of nr or np combination.

Results

The karyotypes of the shrews studied and the expected configurations at meiosis I are shown in Table 2. Four pairs of autosomes (af, bc, jl, tu), as well as the sex chromosomes, were invariant in the karyotypes of all hybrids studied. One pair of acrocentrics (qq) also occurred in all material studied (Table 2).

On the basis of the karyotype of shrew No. 2680, Szałaj *et al.* (1996) decided that the Drnholec race instead of the Stobnica race occurred in the hybrid zone at Plebanka (Fig. 1). Here we wanted to give the precise analysis of this shrew karyotype. This shrew had 24 chromosomes and, among others, chromosomes gand r were acrocentric. The meiotic configurations expected differed clearly according to whether the karyotype included nr or np (Table 2). Analysis of diakinesis figures showed that the first possibility was correct (Fig. 2a). There was no doubt that shrew No. 2680 was a hybrid between the Białowieża and Drnholec races.

A hybrid between the Guzowy Młyn and either the Stobnica or Drnholec races from Wykno (No. 3739) had 23 chromosomes. Again the meiotic configuration revealed that the Drnholec race was involved (Table 2, Fig. 2b, c).

Shrew No. 3758 from the Czarny Piec population had 25 chromosomes and two ik metacentrics diagnostic of the Popielno race. This feature and the presence of only one meiotic complex (Table 1 and 2) suggested that this shrew was a backcross hybrid. The four element ring configuration found at diakinesis of this shrew (Fig. 2d) confirmed that it was a Po × Dn hybrid.

Shrew No. 2541 from the Mycyny population was a hybrid of the Gu \times St or Gu \times Dn type, but had acrocentrics *m* and *n* instead of the *mn* metacentric, and in

No. of shrew, Population	Two variants (<i>nr</i> versus <i>np</i>) of variable part of karyotype	Expected configurations at meiosis I
ropulation	or variable part of Karyotype	at meiosis i
2680	o/ok/ki/ih/hn/nr/r, g/gm/mp/p, qq	str, 5 bi, CH _{VII} , CH _{IV}
Plebanka	o/ok/ki/ih/hn/np/pm/mg/g, rr, qq	str, 6 bi, CH_{IX}
3739	hihi, koko, rg/gm/mn/nr, pp, qq	str, 8 bi, R _{IV}
Wykno	hihi, koko, r/rg/gm/mn/np/p, qq	str, 7 bi, CH_{VI}
3758	hh, kiki, rg/gm/mn/nr, oo, pp, qq	str, 9 bi, R _{IV}
Czarny Piec	hh, kiki, r/rg/gm/mn/np/p, oo, qq	str, 8 bi, CH_{VI}
2541	hihi, koko, m/mg/gr/rn/n, pp, qq	str, 8 bi, CH _{VI}
Mycyny	hihi, koko, m/mg/gr/r, n/np/p, qq	str, 7 bi, CH _{IV} , CH _{III}

Table 2. Partial karyotypes and expected meiotic figures in the hybrids studied. See text for further explanation. Abbreviations: str – sex chromosomes trivalent, bi – bivalent, CH – chain configuration, R – ring configuration, Roman numerals indicate number of elements in complexes.

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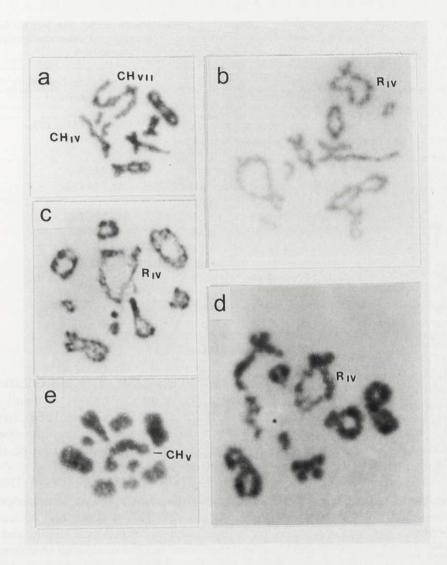


Fig. 2. a – diakinesis of shrew No. 2680, 8 elements, two chains configurations (CH_{VII} and CH_{IV}) are present; b and c – diakinesis of shrew No. 3739, 10 elements, ring of four chromosomes can be seen; d – diakinesis of shrew No. 3758, 11 elements, ring of four chromosomes is present; e – diakinesis of shrew No. 2541, 10 elements; chain of five chromosomes is present.

consequence 24 chromosomes. Two complexes $(CH_{IV} + CH_{III})$ and 7 bivalents would be expected at meiosis if np occurred, and only one (CH_V) complex accompanied by 8 bivalents would indicate the presence of nr (Table 2). A chain-of- five chromosomes was found at diakinesis (Fig. 2e) confirming that this was a Gu × Dn hybrid.

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Table 2. Partial karyotypes and expected meiotic figures in the hybrids studied. See text for further explanation. Abbreviations: str – sex chromosomes trivalent, bi – bivalent, CH – chain configuration, R – ring configuration, Roman numerals indicate number of elements in complexes.

No. of shrew, Population	Two variants (<i>nr</i> versus <i>np</i>) of variable part of karyotype	Expected configurations at meiosis I	
Topulation			
2680	o/ok/ki/ih/hn/nr/r, g/gm/mp/p, qq	str, 5 bi, CH_{VII} , CH_{IV}	
Plebanka	o/ok/ki/ih/hn/np/pm/mg/g, rr, qq	str, 6 bi, CH_{IX}	
3739	hihi, koko, rg/gm/mn/nr, pp, qq	str, 8 bi, R _{IV}	
Wykno	hihi, koko, r/rg/gm/mn/np/p, qq	str, 7 bi, $\mathrm{CH}_{\mathrm{VI}}$	
3758	hh, kiki, rg/gm/mn/nr, oo, pp, qq	str, 9 bi, R_{IV}	
Czarny Piec	hh, kiki, r/rg/gm/mn/np/p, oo, qq	str, 8 bi, $\mathrm{CH}_{\mathrm{VI}}$	
2541	hihi, koko, m/mg/gr/rn/n, pp, qq	str, 8 bi, CH _{VI}	
Mycyny	hihi, koko, m/mg/gr/r, n/np/p, qq	str, 7 bi, CH _{IV} , CH _{III}	

Discussion

Halkka *et al.* (1987) warned of the possibility of mistakes in the identification of chromosomes in the common shrew karyotype, particularly those of small size. Other cytogeneticists have expressed a similar opinion (cf Zima *et al.* 1988). In fact, Halkka *et al.* (1974) and Fredga and Nawrin (1977) labelled chromosome arms m and o conversely, although some features should make identification of these unequivocal when the chromosome preparations are of good quality.

As chromosome arms p and r are also difficult to distinguish from each other we suppose that there are more cases of incorrect identification. Two races described from the vicinity of Moscow are such a case. The first of these had kp and qrmetacentrics and was described from Zvenigorod (Ivanitskaya 1985), and the second one, with combinations kr and pq was found in Chernogolovka (Aniskin and Lukianova 1989). Only the kr and pq metacentrics have recently been confirmed from the region of Moscow (Bulatova *et al.* 2000, Kozlovsky *et al.* 2000). Fusions of pand/or r with arms k, m, n, o and q are characteristic for some other races distributed throughout the whole range of the species (see Zima *et al.* 1996). At least some of these chromosome descriptions could be doubtful.

The present study shows that the Drnholec race has a wider distribution than was described by Wójcik (1993) and Szałaj *et al.* (1996). Although only four individuals have been examined, there is no doubt that in the contact zone between the WEKG and EEKG races in Poland only chromosome nr occurs, both in the southernmost part (Plebanka – the hybrid zone with the Bi race), as well as in the northernmost part (Mycyny – the hybrid zone with the Łg race) (Fig. 3). On the

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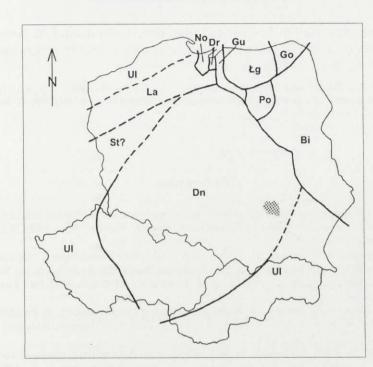


Fig. 3. Proposed distribution of the Drnholec race. Presence of the Stobnica race is doubtful (St?).

other hand, it cannot be excluded that the metacentric np occurs in western Poland. If metacentric np is actually present in Poland, it must have a very small range, limited only to the periphery of the wide range of the metacentric nr (Fig. 3). But, taking the present study into account, occurrence of chromosome np in western Poland needs additional proof.

Establishing that the metacentric nr, instead of np, is present in the hybrid zones is important in the context of our knowledge of the structure of the hybrid zones. Material used for the description of the structure of the hybrid zone near Olsztynek (Fedyk *et al.* 1991, Fedyk 1995, Szałaj *et al.* 1995) needs reexamination. The precise description will be given elsewhere and here we want to correct the basic information about the meiotic complexes that characterize hybrids in the contact zone. In the Dn/Łg hybrid zone (near Olsztynek) two meiotic complexes designated C_1 and C_2 were found in hybrids. The C_1 complex occurs mostly as a ring configuration kh/hi/io/ok (Fedyk *et al.* 1991). Only one pair of chromosomes involved in the C_1 complex is polymorphic (*io*) and with a low frequency of acrocentrics, hence CH_V -forming hybrids o/ok/kh/hi/i are extremely rare (Fedyk *et al.* 1991). The second meiotic complex in the Dn/Łg hybrid zone, designated C_2 , is also a four element ring configuration of rg/gm/mn/nr (Table 1) and not a six element chain, as was described earlier (Fedyk *et al.* 1991). But all the chromosome pairs forming the complex are polymorphic and the frequency of acrocentrics is moderate (Fedyk *et al.* 1991). Hence, the C_2 complex usually occurs as a CH_V or CH_{IV} configuration in hybrids.

Acknowledgements: We are grateful to B. Pochodowicz, M. Sobolewska and A. Wysińska for their help in the field and laboratory work. This paper was sponsored under Grant 6P04C 035 15 from the State Committee for Scientific Research.

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Received 1 September 1999, accepted 4 February 2000.