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Chromosome Complement of *Pomatoschistus marmoratus* and Karyotype Evolution in Sand Gobies (Gobiidae: Gobionellinae)

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Abstract

The family Gobiidae constitutes one of the most diverse groups among teleost fish; some of its species are objects of aquaculture and fisheries. Despite many years of cytogenetic studies, the chromosome complements of many species of gobies remain unknown, and the trends of karyotypic evolution are poorly understood. Here I describe a previously unstudied karyotype of a widespread species from the group of sand gobies (Pomatoschistus marmoratus) and analyze the trends of karyotypic evolution of this group in comparison with other taxonomic groups of gobies from the Atlantic-Mediterranean and Ponto-Caspian regions. For P. marmoratus, the diploid number is 2n=46; it consists of 11 pairs of submetacentric chromosomes, 12 pairs of subtelocentric and telocentric chromosomes (NF=68). A comparative karyological analysis showed that the main trend in the karyotypic evolution of sand gobies differs from that of other groups of gobies and is associated with an increase in the number of chromosomal arms as a result of chromosomal rearrangements leading to a change in the centromere position. The results obtained contribute to understanding the role of chromosomal rearrangements in the evolution of Gobiidae and are of interest for further cytogenetic studies of this large family of fish.

Introduction

Gobies (Gobiidae) are one of the largest families of teleost fish that includes, according to the latest data, more than 1,950 species (Eschmeyer *et al.*, 2021), most of which inhabit the marine and brackish waters of the tropics and subtropics (Nelson *et al.*, 2016). There are also some species of gobies living in freshwater habitats (Freyhof, 2011). Many species of gobies are objects of fisheries and aquaculture (Bell, 1999; Groover *et al.*, 2020; La Mesa *et al.*, 2005).

Among the Atlantic-Mediterranean and Ponto-Caspian fish species, there is a taxonomic group of sand gobies consisting of five genera. Two largest genera are *Knipowitschia* and *Pomatoschistus* which differ in ecology and distribution (McKay & Miller, 1997; Thacker *et al.*, 2019).

The sand gobies are characterized by a number of common cytogenetical features, namely: in most species, the diploid number of chromosomes (2n) is 46; the nucleolar organizer regions are located in the terminal-centromeric zone on the short arm in one submetacentric pair of chromosomes; they have similar values of the DNA amount per cell (Caputo, 1998; Rampin et al., 2011). Molecular phylogenetic data calibrated using fossils indicate the origin of the sand goby clade near the Eocene-Oligocene boundary (about 33.0 Mya) (Thacker et al., 2019). The relatively long evolutionary history of this group of gobies (Huyse et al., 2004; Malavasi et al., 2012) was reflected in their karyotypic diversity. Among the karyologically analyzed species of the sand gobies, the number of chromosomes varies from 2n=32 to 2n=46, while the fundamental arm number (NF) varies from NF=50 to NF=70. Nevertheless,

the chromosome complement of a number of sand goby species remains unknown, and karyotypic evolution remains poorly understood.

In the present work, I describe the chromosome complement of a previously unstudied species (*Pomatoschistus marmoratus*) and a comparative karyological analysis of a group of sand gobies. The results obtained showed that the main trend of karyotypic evolution of this group differs from that of other taxonomic groups of gobies and is associated with an increase in the proportion of two-armed chromosomes (metacentric and submetacentric) as a result of rearrangements leading to a change in the centromere position. This work contributes to the expansion of knowledge of the mechanisms for the formation of karyotypic diversity and creates opportunities for further cytogenetic studies of the family Gobiidae.

Materials and Methods

The material for karyological analysis was collected in the Black Sea, near the coast of the Bugaz Spit (45°04'15.5"N, 37°00'48.0"E). A total of 12 specimens of *Pomatoschistus marmoratus* (Risso, 1810) were caught. Most of the specimens were adult males with a total length (*TL*) of 53–58 mm (Figure 1). Mitotic activity sufficient for chromosome analysis was observed in 5 specimens (4 males and 1 female). A total of 67 metaphase plates were studied.

Chromosome preparations were obtained from the anterior part of the kidney according to previously published methods (Blanco *et al.*, 2012; Ojima & Kurishita, 1980) with the initial treatment of live fish with colchicine (injection of 0.03% solution into the spinal muscle). The anterior kidney tissue was incubated in 75 mM KCl (hypotonic solution) for 22 min at 27°C and fixed in 96% ethanol mixed with glacial acetic acid (3:1 ratio). Chromosome preparations made using standard air-drying techniques were stained with 5% Giemsa solution in phosphate buffer at pH 6.8 for 7 min.

Analysis of mitotic chromosomes was performed using the KaryoType software (Altinordu *et al.*, 2016). Chromosomes were classified according to Levan et al. (1964): metacentric (m), submetacentric (sm), subtelocentric (st) and telocentric (t). To determine the fundamental arm number (NF), chromosomes of the m and sm groups were considered two-armed and those of the st/t group one-armed. The regression between the proportion of two-armed chromosomes and diploid chromosome number and the Spearman correlation were calculated. For statistical analysis of the results, the Excel 2016 program was used.

Results and Discussion

Karyological studies of *Pomatoschistus marmoratus* from the Black Sea near the coast of the Bugaz Spit showed the chromosome complement containing 2n=46. The karyotype consists of 11 pairs of submetacentric and 12 pairs of subtelocentric and telocentric chromosomes; the fundamental arm number NF=68 (Figure 2). No sex chromosomes were found in the studied chromosome complements. A characteristic feature of the *P. marmoratus* karyotype is the presence of pairs of large submetacentric and



Figure 1. A male of *Pomatoschistus marmoratus*. Scale bar – 5 mm.



Figure 2. Karyotype of Pomatoschistus marmoratus from the Black Sea (2n=46). Scale bar – 10 µm.

TRJFAS20729

telocentric chromosomes (Figures 2, 3). A marker pair of large telocentric chromosomes has also been previously found in the karyotypes of other sand goby species: *Pomatoschistus bathi, P. lozanoi, P. minutus* and *P. norvegicus* (Boltachev *et al.,* 2016; Webb, 1986).

The diploid karyotype of most of the studied sand gobies consists of 46 chromosomes (Table 1). The modal number of chromosomes is 2n=46 both for the goby groups of Atlantic-Mediterranean, Ponto-Caspian and sand gobies (Figure 4) and, apparently, for the entire suborder Gobioidei (da Silva *et al.*, 2021; Prazdnikov, 2013). Moreover, from a cytogenetic point of view, the diploid karyotype consisting of 46 one-armed chromosomes is the most primitive and should be considered as the basic/ancestral karyotype of goby fish.

An analysis of karyological data (Arai, 2011; Galvão *et al.*, 2011; Prazdnikov, 2013) indicated that the evolution of the karyotype in goby fish occurred at different rates and in different directions, which led to a wide chromosomal divergence (2n=29–56, NF=38–96).

For the Atlantic-Mediterranean and Ponto-Caspian groups of goby fish, the proportion of two-armed chromosomes in the karyotype varies widely from 0% to 59%. The regression between the proportion of two-armed chromosomes in the karyotype and the diploid number is y=-2.135x+103.77 (R^2 =0.63), and the Spearman correlation is R_s=-0.604 (Figure 5a). For the group of sand gobies, the proportion of two-armed

chromosomes in the karyotype varies from 13% to 65%. The regression between the proportion of two-armed chromosomes and the diploid number is y=-1.1509x+90.689 (R²=0.08); R_s=0 (Figure 5b). The absence of a statistically significant association between the two variables (2n and proportion of m/sm chromosomes) in sand gobies is apparently due to chromosomal rearrangements that affected the trends of karyotypic evolution in this group.

The main trend in the karyotypic evolution of most groups of gobies from the Atlantic-Mediterranean and Ponto-Caspian regions is the result of the rearrangements of the centric fusion type (Robertsonian translocations) leading to a decrease in the number of chromosomes (without changing the fundamental arm number) (Figure 6a). Another direction of karyotype transformation associated with an increase in the number of chromosomes (as a result of centric fission) and chromosomal arms (mainly as a result of pericentric inversion) is less pronounced.

In contrast, the main trend of karyotypic evolution for the group of sand gobies is associated with an increase in the number of chromosomal arms (without changing the number of chromosomes) as a result of rearrangements leading to a change in the centromere position, mainly due to pericentric inversion (Figure 6b). As a result, the karyotypes of *Economidichthys pygmaeus* and some populations of *Pomatoschistus*



Figure 3. Ideogram of Pomatoschistus marmoratus (n=23). Scale bar – 3 µm.

Species	2n	Karyotype formula	NF	References
Economidichthys pygmaeus	46	24sm+22st	70	Rampin et al., 2011
Pomatoschistus bathi	44	6sm+38st/t	50	Boltachev et al., 2016
Pomatoschistus flavescens	46	6m/sm+40t	52	Klinkhardt, 1992
Pomatoschistus lozanoi	37	3m+12sm+10st+12t	52	Webb, 1980
Pomatoschistus marmoratus	46	22sm+24st/t	68	Present study
Pomatoschistus microps	46	30sm+16st	76	Webb, 1986
	46	4m+16sm+20st+6t	66	Klinkhardt, 1989
Pomatoschistus minutus	46	4m+16sm+16st+10t	66	Klinkhardt, 1989
	46	18sm+18st+10t	64	Klinkhardt, 1992
	46	6sm+24st+16t	52	Webb, 1980
Pomatoschistus norvegicus	32	10m+10sm+8st+4t	52	Webb, 1980
Pomatoschistus pictus	46	22m/sm+12st+12t	68	Klinkhardt, 1992

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Table 1. C	ytogenetic	data for	species	of sand	gobies



Figure 4. Histogram of the distribution of the diploid chromosome number: (a) for the Atlantic-Mediterranean and Ponto-Caspian groups of gobies, (b) for the group of sand gobies. An odd number of chromosomes is associated with either heterozygous karyotype variants or Y-autosomal translocation.



Figure 5. Scatter-plot of a diploid chromosome number and proportion of metacentric/submetacentric chromosomes (m/sm) with overall regression line: (a) for the Atlantic-Mediterranean and Ponto-Caspian groups of gobies, (b) for the group of sand gobies. The diameter and color of a circle indicate the number of species from 1 to 15.

microps consist exclusively of submetacentric and subtelocentric chromosomes (Table 1) (Rampin et al., 2011; Webb, 1986). An additional direction of karyotype transformations of sand gobies is associated with a decrease in the number of chromosomes due to Robertsonian translocations and an increase in the number of chromosome arms due to pericentric inversions relative to the basic chromosome complement (Figure 6b). This direction is typical for the karyotypes of Pomatoschistus lozanoi and P. norvegicus (Table 1). Interestingly, the P. lozanoi karyotype is heterozygous for the Robertsonian translocation (2n=37), and the appearance of an unpaired metacentric element is not associated with sex chromosomes (Webb, 1980). It should be noted that changes in the position of the centromere in the chromosome complements of sand gobies are probably caused not only by pericentric inversions, but also by the formation of neocentromeres which can be observed in the karyotypes of various groups of animals (Schubert, 2018; Sobita & Bhagirath, 2006).

Among the karvologically studied species of sand gobies, two species, Pomatoschistus microps and P. minutus, are characterized by the interpopulation differentiation of karyotypes (Table 1). It is possible that the karyotypic diversity of sand gobies may be increasing as a result of hybridization. To date, there are known cases of hybridization in sympatric zones between P. microps and P. marmoratus (Berrebi & Trébuchon, 2020), as well as between P. lozanoi and P. norvegicus, and P. lozanoi and P. minutus (Wallis & Beardmore, 1980; Webb, 1980). Thus, it can be expected that in the course of further cytogenetic studies both homozygotes and heterozygotes for various chromosomal rearrangements that affect the fertility of sand goby hybrids shall be found in sympatric zones.

Analysis of the karyotypic evolution of different taxa of fish shows that the probability of a purely random fixation of chromosomal rearrangements without the participation of natural selection is very small, which is evidence in favor of the adaptive significance of such rearrangements (Cayuela *et al.*,

2020; Kirpichnikov, 1987; Kirubakaran et al., 2016; Martinez et al., 2015). It has been earlier suggested that the karyological differentiation in sand gobies of the genus Pomatoschistus may have an adaptive value (Webb, 1980). The relatively ancient origin of Pomatoschistus about 18.3 Mya and major tectonic events in the Mediterranean basin contributed to the diversification of sand gobies (Thacker et al., 2019). A series of complex hydrographic and geological events associated with the isolation of the Tethys and Paratethys seas in the Miocene with subsequent epochs of regression and elevation of the water levels, as well as fluctuations in salinity, led first to the isolation of the goby lineages and then to their adaptive radiation (Huyse et al., 2004; Malavasi et al., 2012; Thacker, 2015). Probably, all these events influenced the ecology and distribution of sand gobies and led to the formation of karyotypes composed of a large number of twoarmed chromosomes.

The specifics of the complex history of the Mediterranean basin also influenced the karyotypic diversification of other fish taxa with the formation of intrapopulational and interpopulational chromosomal polymorphism (Prazdnikov, 2016; Vitturi *et al.*, 1986; Vitturi & Lafargue, 1992).

Conclusion

The diploid karyotype of Pomatoschistus marmoratus, one of the widespread sand goby species, consists of 46 chromosomes with a high proportion of submetacentric elements. A comparative karyological analysis showed that the trends of chromosomal evolution in the sand goby group differ from those of other taxonomic groups of gobies. In the karyotypes of the proportion of sand gobies, two-armed chromosomes increases without changes in the basic number of chromosomes (2n=46), mainly due to pericentric inversions. The ancient origin of the sand goby group together with a series of hydrographic and geological events in the region of their habitat, as well as differences in ecology and distribution, provide evidence in favor of the potentially adaptive value of



Figure 6. The trends of karyotypic evolution (a) for the Atlantic-Mediterranean and Ponto-Caspian groups of gobies, (b) for the group of sand gobies. The red thick arrow shows the most probable main trend in karyotypic evolution. The lower rectangle shows the ancestral karyotype. 2n - number of chromosomes in a diploid complement, NF – fundamental arm number, fu – centric fusions, fiss – centric fissions, per inv – pericentric inversions.

karyological differentiation. Taking into account that some sand gobies are characterized by a disjunct distribution with overlapping geographic ranges between species, further cytogenetic studies are likely to reveal an even greater karyotypic diversity in this group of fish.

Ethical Statement

Not applicable

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Conflict of Interest

The author declares no conflict of interest.

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