

Chromosomal complements of some Atlantic Blennioidei and Gobioidae species (Perciformes)

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Abstract

A remarkable degree of chromosomal conservatism ($2n=48$, $FN=48$) has been identified in several families of Perciformes. However, some families exhibit greater karyotypic diversity, although there is still scant information on the Atlantic species. In addition to a review of karyotypic data available for representatives of the suborders Blennioidei and Gobioidae, we have performed chromosomal analyses on Atlantic species of the families Blenniidae, *Ophioblennius trinitatis* Miranda-Ribeiro, 1919 ($2n=46$; $FN=64$) and *Scartella cristata* (Linnaeus, 1758) ($2n=48$; $FN=50$), Labrisomidae, *Labrisomus nuchipinnis* (Quoy & Gaimard, 1824) ($2n=48$; $FN=50$) and Gobiidae, *Bathygobius soporator* (Valenciennes, 1837) ($2n=48$; $FN=56$). Besides variations in chromosome number and karyotype formulas, Ag-NOR sites, albeit unique, were located in different positions and/or chromosome pairs for the species analyzed. On the other hand, the heterochromatic pattern was more conservative, distributed predominantly in the centromeric/pericentromeric regions of the four species. Data already available for Gobiidae, Blenniidae and Labrisomidae show greater intra- and interspecific karyotypic diversification when compared to other groups of Perciformes, where higher uniformity is found for various chromosome characteristics. Evolutionary dynamism displayed by these two families is likely associated with population fractionation resulting from unique biological characteristics, such as lower mobility and/or specific environmental requirements.

Keywords

Chromosomal evolution, marine fish, Blenniidae, Gobiidae, Labrisomidae

Introduction

Although karyotypic characteristics for some families of marine fish are already known, information on groups of Perciformes is still significantly disproportionate. Among these, suborders Blennioidei and Gobioidae stand out because of the large number of species they represent.

Suborders Gobioidae, with 2,121 species, and Blennioidei with 732 species, are spread throughout the tropical zone, typically represented by small specimens with low mobility and the ability to withstand changes in temperature and salinity (Nelson 2006).

Species of Blennioidei and Gobioidae investigated (e.g. Cataudela et al. 1973; Garcia et al. 1987; Ene 2003) have shown sufficient chromosomal peculiarities for species discrimination and understanding of their evolutionary aspects. In some families, such as Blenniidae, Labrisomidae and Gobiidae, sharing cryptic morphological characteristics combined with poor knowledge of the biological characteristics for many species, contributes to the relative taxonomic inaccuracy of this group. As such, cytotaxonomic markers (Garcia et al. 1987; Caputo 1998; Caputo et al. 2001) and phylogenetic analyses based on molecular data (Wang et al. 2001; Thacker 2003; Gysels et al. 2004; Almada et al. 2005) have been increasingly used when assessing their kinship relations. Indeed, it has been suggested that phylogenetic analyses combine molecular and morphological data (Thacker 2003), as well as cytogenetic information. However, in light of the diversity in these groups, solid chromosome data are not yet sufficiently available, with only 7.5% of Blenniidae species and 4.5% of Gobioidae was karyotyped (Table 1). Despite the scarcity of data, a high degree of chromosomal polymorphism has been characterized among Gobiidae, primarily Robertsonian rearrangements (Caputo et al. 1999, Ene 2003), along with others such as tandem fusions and pericentric inversions (Giles et al. 1985; Thode et al. 1985; Amores et al. 1990).

The present study focuses on the karyotypic characterization of some Atlantic species of the families Blenniidae, *Ophioblennius trinitatis* Miranda-Ribeiro, 1919 and *Scartella cristata* (Linnaeus, 1758), Labrisomidae, *Labrisomus nuchipinnis* (Quoy & Gaimard, 1824) and Gobiidae, *Bathygobius soporator* (Valenciennes, 1837), through conventional chromosomal analysis, characterization of nucleolar organizer regions (Ag-NORs) and the distribution pattern of C-positive heterochromatin (C-banding) in chromosomes, discussing evolutionary aspects.

Material and methods

A total of 25 specimens of *Ophioblennius trinitatis* (7♂, 4♀ and 14 indeterminate), 11 specimens of *Scartella cristata* (4♂, 5♀ and 2 indeterminate), 13 specimens of *Labrisomus nuchipinnis* (4♂, 4♀ and 5 indeterminate) and 12 specimens of *Bathygobius soporator*, (5♂, 5♀ and 2 indeterminate) were used for chromosome analysis. *Ophioblennius trinitatis* specimens came from the coast of Rio Grande do Norte

Table 1. Cytogenetic data for Blennioidei and Gobioidaei (Perciformes).

Suborder/ Family	Species	2n	Karyotype formula	FN	References
Blennioidei					
Blenniidae	<i>Aidablennius sphyinx</i>	48	4m+4sm+40a	56	Cano et al. (1982)
	<i>A. sphyinx</i>	48	2st+46a	50	Cataudella and Civitelli (1975)
	<i>Atrosalarias fuscus</i>	48	48a	48	Arai and Shiotsuki (1973)
	<i>Blennius ocellaris</i>	48	2m+2st+44a	52	Vitturi et al. (1986)
	<i>B. ponticus</i>	48	16sm+10st+22a	74	Garcia et al. (1987)
	<i>B. yatabei</i>	48	6sm+12st+30a	66	Arai and Shiotsuki (1974)
	<i>Coryphoblennius galerita</i>	48	2m+12sm+34a	62	Garcia et al. (1973)
	<i>Dasson trossulus</i>	40	8m+32st/a	48	Arai and Shiotsuki (1974)
	<i>Istiblennius enoshimae</i>	48	2m+46a	50	Arai and Shiotsuki (1973)
	<i>I. lineatus</i>	48	48st/a	48	Arai and Shiotsuki (1974)
	<i>Lipophrys canevai</i>	48	8st+40a	56	Cataudella and Civitelli (1975)
	<i>L. pholis</i>	46	8m+8sm+30a	62	Garcia et al. (1987)
	<i>L. trigloides</i>	46	4m+4sm+10st+28a	64	Cano et al. (1982)
	<i>L. trigloides</i>	48	2m+6sm+18st+22a	74	Cataudella and Civitelli (1975)
	<i>L. trigloides</i>	48	2m+22sm+2st+22a	74	Garcia et al. (1987)
	<i>L. trigloides</i>	48	2m+6sm+18st+22a	74	Vitturi et al. (1986)
	<i>Omobranchus elegans</i>	42	10m+2sm+6st+24a	60	Arai and Shiotsuki (1974)
	<i>O. punctatus</i>	44	4m+40a	48	Arai (1984)
	<i>Ophioblennius trinitatis</i>	46	6m+12st+28a	64	Present study
	<i>Parablennius incognitus</i> (= <i>Blennius incognitus</i>)	48	4st+44a	52	Cano et al. (1982)
	<i>P. pilicornis</i> (= <i>Blennius pilicornis</i>)	48	8st+40a	56	Catalano et al. (1985)
	<i>P. gattorugine</i>	48	2m+4sm+42a	54	Vitturi et al. (1986)
	<i>P. pilicornis</i>	48	48a	48	Brum et al. (1992)
	<i>P. sanguinolentus</i>	48	12st+36a	60	Cataudella et al. (1973)
	<i>P. sanguinolentus</i>	48	20sm+10st+18a	78	Garcia et al. (1987)
	<i>P. tentacularis</i>	48	48st/a	48	Vasil'ev (1985)
	<i>P. tentacularis</i>	48	1st+47a	49	Carbone et al. (1987)
	<i>P. tentacularis</i>	47	1sm+46a	48	Carbone et al. (1987)
	<i>Salarias fluviatilis</i>	48	48st/a	48	Cataudella and Civitelli (1975)
	<i>S. pavo</i>	48	8st+40a	56	Cataudella et al. (1973)
	<i>S. pavo</i>	48	16sm+14st+18a	78	Garcia et al. (1987)
	<i>S. pavo</i>	48	2st+46a	50	Vasil'ev (1980)
	<i>Salarias faciatus</i>	48	48a	48	Arai and Shiotsuki (1973)
	<i>S. luctuosus</i>	48	48st/a	48	Arai and Shiotsuki (1974)
	<i>Scartella cristata</i> (= <i>Blennius cristatus</i>)	48	2st+46a	50	Vitturi et al. (1986)
	<i>S. cristata</i>	48	2sm+46st/a	50	Brum et al. (1995)
	<i>S. cristata</i>	48	4st+44a	52	Present study
Gobioidaei					
Clinidae	<i>Clinithractus argentatus</i>	48	2st+46a	50	Vitturi et al. (1986)
Labrisomidae	<i>Labrisomus nuchipinnis</i>	48	2sm+46a	50	Affonso (2000)
	<i>L. nuchipinnis</i>	48	2st+46a	50	Present study

Suborder/ Family	Species	2n	Karyotype formula	FN	References
Eleotridae	<i>Dormitator latifrons</i>	46	44m/sm+2st/a	90	Uribe-Alcocer et al. (1983)
	<i>D. maculatus</i>	46	34m/sm+12st/a	80	Maldonado-Monroy et al. (1985)
	<i>D. maculatus</i>	46	40m/sm+6st/a	86	Molina (2005)
	<i>D. maculatus</i>	46	14m+28sm+2st+2a(♀) 13m+28sm+3st+2a(♂)	90	Oliveira and Almeida-Toledo (2006)
	<i>Eleotrioides strigatus</i>	44	2m+42st/a	46	Arai and Sawada (1974)
	<i>Eleotris acanthopomus</i>	46	46st/a	46	Arai and Sawada (1974)
	<i>E. picta</i>	52	52a	52	Uribe-Alcocer and Diaz-James (1996)
	<i>E. pisonis</i>	46	2m/sm+42st/a	46	Uribe-Alcocer and Diaz-James (1996)
	<i>E. pisonis</i>	46	46a	46	Rocon-Strange (1992)
	<i>E. pisonis</i>	46	46a	46	Molina (2005)
	<i>E. muralis</i>	46	46a	46	Khuda-Bukhsh and Nayak (1990)
	<i>Mogurnda mogurnda</i>	46	6sm+40st/a	52	Arai et al. (1974)
	<i>M. obscura</i>	62	-	-	Nogusa (1960)
	<i>Ophiocara porocephala</i>	48	48a	48	Arai and Fujiki (1979)
	<i>Oxyeleotris marmorata</i>	46	2m+2sm+42a	50	Arai and Fujiki (1979)
Gobiidae	<i>Aboma latipes</i>	40	40a	40	Arai and Sawada (1974)
	<i>Acanthogobius flavimanus</i>	44	44st/a	44	Arai and Sawada (1974)
	<i>A. flavimanus</i>	44	36st+8a	80	Arai and Kobayashi (1973)
	<i>A. flavimanus</i>	44	10m/sm/st+34a	54	Arai and Sawada (1975)
	<i>Acentrogobius pflaumi</i>	50	48m/sm+2st/a	98	Nogusa (1960)
	<i>Amblygobius albimaculatus</i>	44	2m+42st/a	46	Nishikawa et al. (1974)
	<i>Aphia minuta</i>	44	44a	44	Caputo et al. (1999)
	<i>A. minuta</i>	43	42a+1st	42	Caputo et al. (1999)
	<i>A. minuta</i>	42	1m+1st+40a	44	Caputo et al. (1999)
	<i>A. minuta</i>	42	1M+1m+40a	44	Caputo et al. (1999)
	<i>A. minuta</i>	41	2M+1st+38a	44	Caputo et al. (1999)
	<i>Apocryptes bato</i>	46	24m+10sm+12a	80	Nayak and Khuda-Bukhsh (1987)
	<i>A. lanceolatus</i>	38	14m+22sm+2st	76	Nayak and Khuda-Bukhsh (1987)
	<i>Awaous grammepomus</i>	46	46st/a	46	Khuda-Bukhsh and Barat (1987)
	<i>A. tajasica</i>	46	46a	46	Strange and Passamani (1986)
	<i>Bathygobius fuscus</i>	48	48a	48	Arai and Sawada (1975)
	<i>B. saporator</i>	48	2m+46a	50	Brum et al. (1996)
	<i>B. saporator</i>	48	2m/sm+46a	50	Cipriano et al. (2002)
	<i>B. saporator</i>	48	2m+6st+40a	56	Present study
	<i>B. stellatus</i>	46	2st+44a	48	Vasil'ev (1985)
	<i>B. stellatus</i>	47	1sm+2st+43a	49	Vasil'ev (1985)
	<i>Boleophthalmus boddarty</i>	46	46m/sm	92	Subrahmanyam (1969)
	<i>B. glaucus</i>	46	12m+20sm+2st+12a	80	Manna and Prasad (1974)
	<i>B. pectinirostris</i>	46	46st/a	46	Arai and Sawada (1975)
	<i>Bostrichthys sinensis</i>	48	4m/sm+44a	52	Arai et al. (1974)
	<i>Chaenogobius annularis</i>	44	18sm+26st/a	62	Arai and Sawada (1975)
	<i>C. annularis</i>	44	36m/sm+8a	80	Arai et al. (1974)
	<i>C. annularis</i>	44	44a	44	Nogusa (1960)
	<i>C. castaneus</i>	44	36m/sm/st+8a	80	Nishikawa et al. (1974)
	<i>C. isaza</i>	44	12sm+32st/a	56	Arai and Sawada (1975)
	<i>C. urotaenia</i>	44	-	-	Nogusa (1960)

Suborder/ Family	Species	2n	Karyotype formula	FN	References
	<i>C. urotaenia</i>	42	14sm+28a	56	Yamada (1967)
	<i>Chasmichthys dolichognatus</i>	44	44st/a	44	Arai and Sawada (1975)
	<i>C. gulosus</i>	44	44st/a	44	Arai and Sawada (1975)
	<i>C. gulosus</i>	44	16m/sm/st+28a	60	Nishikawa et al. (1974)
	<i>Ctenogobius criniger</i>	50	34m/sm+6st+10a	90	Arai and Sawada (1974)
	<i>Gillichthys mirabilis</i>	44	12sm+32a	56	Chen and Ebeling (1971)
	<i>G. seta</i>	44	6m+14sm+24a	64	Chen and Ebeling (1971)
	<i>Glossogobius fasciatopunctatus</i>	44	10m+28sm+2st+4a	84	Fei and Tao (1987)
	<i>G. giuris</i>	46	46a	46	Rishi and Singh (1982)
	<i>Gobiodon citrinus</i>	44	2m+42st/a	46	Arai and Sawada (1974)
	<i>G. citrinus</i>	43	1m+42st/a	44	Arai and Sawada (1974)
	<i>G. quinquestrigatus</i>	44	44a	44	Arai and Fujiki (1979)
	<i>G. rivulatus</i>	44	44a	44	Arai and Fujiki (1979)
	<i>Gobioides rubicundus</i>	46	2m+26sm+10st+8a	84	Manna and Prasad (1974)
	<i>Gobionellus shufeldti</i>	48	48a (♀)	48	Pezold (1984)
	<i>G. shufeldti</i>	47	46a+1m (♂)	48	Pezold (1984)
	<i>Gobiosoma macrodon</i>	38	38a	38	Musammil (1974)
	<i>G. zebrella</i>	38	38a	38	Musammil (1974)
	<i>Gobius abei</i>	46	-	-	Nogusa (1960)
	<i>G. buccichi</i>	44	2sm+42a	46	Thode and Alvarez (1983)
	<i>G. cobitis</i>	46	46a	46	Caputo et al. (1997)
	<i>G. cruentatus</i>	46	2st+44a	48	Thode and Alvarez (1983)
	<i>G. fallax</i>	38	8m/sm+30a	46	Thode et al. (1988)
	<i>G. fallax</i>	39	7m/sm+32a	46	Thode et al. (1988)
	<i>G. fallax</i>	40	6m/sm+34a	46	Thode et al. (1988)
	<i>G. fallax</i>	40	7m/sm+33a	47	Thode et al. (1988)
	<i>G. fallax</i>	41	5m/sm+36a	46	Thode et al. (1988)
	<i>G. fallax</i>	42	4m/sm+38a	46	Thode et al. (1988)
	<i>G. fallax</i>	43	3m/sm+40a	46	Thode et al. (1988)
	<i>G. niger</i>	52	2m+4sm+16st+30a	74	Vitturi and Catalano (1989)
	<i>G. niger</i>	51	3m+4sm+16st+28a	74	Caputo et al. (1997)
	<i>G. niger</i>	50	4m+4sm+16st+26a	74	Caputo et al. (1997)
	<i>G. niger</i>	49	5m+4sm+16st+24a	74	Caputo et al. (1997)
	<i>G. paganellus</i>	48	2sm+46a	50	Caputo et al. (1997)
	<i>G. similis</i>	44	?		Nogusa (1960)
	<i>Gobiusculus flavescens</i>	46	6m/sm+40a	52	Klinkhardt (1992)
	<i>Luciogobius grandis</i>	44	?		Arai (1981)
	<i>L. guttatus</i>	44	?		Arai and Kobayashi (1973)
	<i>Mesogobius batrachocephalus</i>	30	16m+14a	46	Ivanov (1975)
	<i>Neogobius cephalarges</i>	46	46a	46	Vasil'ev (1985)
	<i>N. constructor</i>	42	4m/sm+38a	46	Vasil'ev and Vasil'yeva (1994)
	<i>N. cyrius</i>	36	structural polymorphism		Vasil'ev and Vasil'yeva (1994)
	<i>N. fluviatilis</i>	46	46a	46	Vasil'ev (1985)
	<i>N. eurycephalus</i>	32	12m+2sm+18a	46	Ene (2003)

Suborder/ Family	Species	2n	Karyotype formula	FN	References
	<i>N. eurycephalus</i>	31	13m+2sm+16a	46	Ene (2003)
	<i>N. eurycephalus</i>	30	14m+2sm+14a	46	Ene (2003)
	<i>N. gymnotrachelus</i>	46	46a	46	Vasil'ev and Grigoryan (1992)
	<i>N. kessleri</i>	46	46a	46	Vasil'ev (1985)
	<i>N. melanostomus</i>	46	46a	46	Vasil'ev (1985)
	<i>N. rhodionovi</i>	46	46a	46	Vasil'ev and Vasil'yeva (1994)
	<i>Odontamblyops rubicundus</i>	46	4m+16sm+26st/a	66	Arai and Sawada (1975)
	<i>Padogobius martensi</i>	46	1m+3sm+2st+40a	52	Cataudella et al. (1973)
	<i>Parioglossus raoi</i>	46	46st/a	46	Webb (1986)
	<i>Periophthalmus cantonensis</i>	46	18m+12sm+16st/a	76	Arai and Sawada (1975)
	<i>Pomatoschistus lozanoi</i>	37	3m+12sm+10st+12a	62	Webb (1980)
	<i>P. microps</i>	46	4m+16sm+20st+6a	86	Klinkhardt (1989)
	<i>P. minutus</i>	46	4m+16sm+16st+10a	82	Klinkhardt (1989)
	<i>P. minutus</i>	46	18sm+18st+10a	82	Klinkhardt (1992)
	<i>P. norvegicus</i>	32	10m+10sm+8st+4a	60	Webb (1980)
	<i>P. pictus</i>	46	22m/sm+12st+12a	80	Klinkhardt (1992)
	<i>Proterorhinus marmoratus</i>	46	46a	46	Rab (1985)
	<i>Pterogobius elapoides</i>	44	14sm+30st	88	Arai and Kobayashi (1973)
	<i>P. zonoleucus</i>	44	14sm+30st	88	Arai and Sawada (1975)
	<i>Quietula guaymasiae</i>	42	6m+4sm+32a	52	Cook (1978)
	<i>Q. y-cauda</i>	42	42a	42	Cook (1978)
	<i>Rhinogobius brunneus</i>	44	44a	44	Nishikawa et al. (1974)
	<i>R. flumineus</i>	44	44a	44	Arai and Kobayashi (1973)
	<i>R. giurinus</i>	44	44a	44	Nishikawa et al. (1974)
	<i>Rhodoniichthys laevis</i>	42	16m/sm+26st	84	Arai et al. (1974)
	<i>Sicyopterus japonicus</i>	44	10m+10sm+24a	64	Arai and Fujiki (1979)
	<i>Synechogobius hasta</i>	44	2m+42st/a	46	Arai and Sawada (1975)
	<i>Tridentiger obscurus</i>	44	10m/sm+34a	54	Arai et al. (1974)
	<i>T. trigonocephalus</i>	44	28m/sm/st+16a	72	Arai et al. (1973)
	<i>T. trigonocephalus</i>	46	16sm+6st+24a	68	Fei and Tao (1987)
	<i>Trypauchen vagina</i>	46	12m+6sm+10st+18a	74	Khuda-Bukhsh (1978)
	<i>Tukugobius flumineus</i>	44	44a	44	Nadamitsu (1974)
	<i>Zosterisessor ophiocephalus</i> (= <i>Gobius ophiocephalus</i>)	46	46a	46	Vasil'ev (1985)
	<i>Zosterisessor ophiocephalus</i> (= <i>Gobius ophiocephalus</i>)	45	1st+45a	47	Vasil'ev (1985)
	<i>Zosterisessor ophiocephalus</i>	46	2m/sm+44a	48	Caputo et al. (1996)

(5°13'1.73"S; 35°9'57.85"W), northeastern Brazil (n=1), and the Saint Peter and Saint Paul (n=8) (00°55'02"N; 29°20'42"W) and Fernando de Noronha (n=16) (3°52'11"S; 32°26'13"W) archipelagos. The remaining specimens were collected on the coast of Rio Grande do Norte. Individuals were previously submitted to mitotic stimulation with compound attenuated antigens, for 24 to 48 hours (Molina 2001, Molina et al. 2010), anesthetized with clove oil (Eugenol) and sacrificed for the removal of anterior kidney fragments. Sexing of specimens was performed by macroscopic and microscopic examination of the gonads. Chromosome preparations were obtained from kidney

cells (Gold et al. 1990). Nucleolar organizer regions (NORs) were identified by stain with silver nitrate - Ag-NORs (Howell and Black 1980) and C-positive heterochromatin sites through C-banding (Sumner 1972).

Metaphase preparations were examined and photographed on an Olympus BX50 photomicroscope, using an Olympus DP70 digital camera system. Chromosomes were classified according to the position of the centromere in metacentrics (m), submetacentrics (sm), subtelocentrics (st) and acrocentrics (a) (Levan et al. 1964) and organized in order of decreasing size. The chromosome formula and FN (fundamental number or number of chromosomal arms) were established for each species, considering acrocentric chromosomes with a single arm and the remaining chromosomes exhibiting two arms.

Results

Cytogenetic analyses of Blennioidei species (Blennioidei)

Ophioblennius trinitatis showed $2n=46$, with a chromosome formula equal to $6m+12st+28a$ (FN=64), irrespective of sex. Although chromosomes showed a gradual decline in size, the smallest acrocentric pairs corresponded to approximately one-third of the largest metacentric pairs. Nucleolar organizer regions are located in the terminal portions of the short arm on pair 9, the smallest subtelocentric pair. C-positive heterochromatin is discretely located in the centromeric/pericentromeric region of the chromosomes (Fig. 1a, b).

Scartella cristata showed $2n=48$ chromosomes, with a chromosome formula equal to $4st+44a$ (FN=52). The karyotype also displays a gradual reduction in chromosome size. However, the largest chromosome pair exhibits only double the size in relation to the smallest karyotype pair. Ribosomal sites are located on the terminal portions of the short arms on chromosome pair 1. C-positive heterochromatin is also reduced and located in the centromeric regions of chromosomes (Fig. 1c, d).

Cytogenetic analyses of Labrisomidae and Gobiidae species (Gobioidaei)

Labrisomus nuchipinnis (Labrisomidae) showed $2n=48$ chromosomes with a chromosome formula of $2st+46a$ (FN=50), showing relatively more differentiated size between the largest and smallest chromosomes of the karyotype. Nucleolar organizer regions are in the terminal portions of the long arms on pair 2, corresponding to the largest pair of acrocentric chromosomes. C-positive heterochromatin was showed in the centromeric/pericentromeric region of all chromosome pairs, in relatively conspicuous blocks (Fig. 1e, f).

Bathygobius soporator (Gobiidae) also displayed the karyotype composed of $2n=48$ chromosomes, but with the chromosome formula distinct from that of *L. nuchipinnis*, specifically, $2m+6st+40a$ (FN=56). Size difference between the largest and smallest

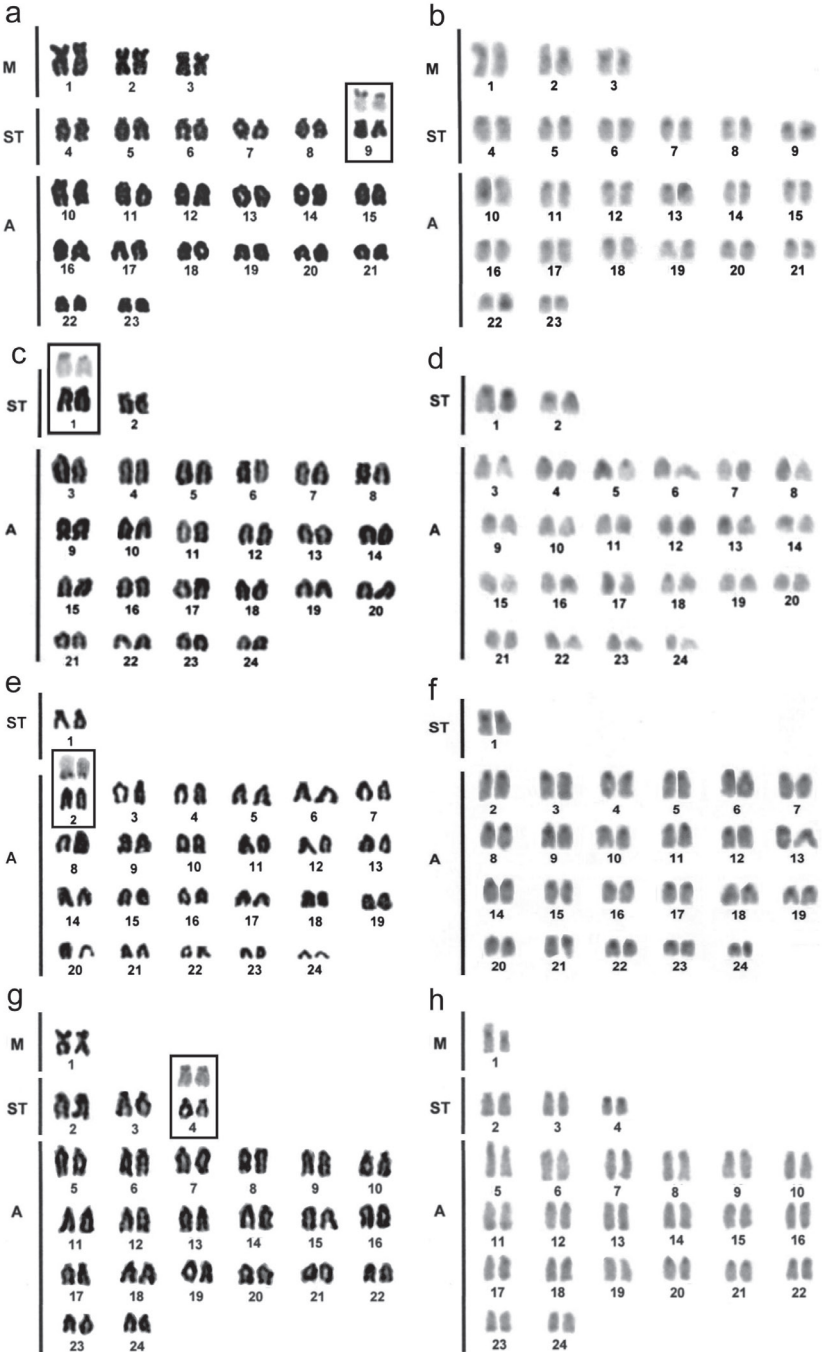


Figure 1. Karyotypes under Giemsa staining **a, c, e, g** and C-banding **b, d, f, h** of *Ophioblennius trinitatis*; **a, b** *Scartella cristata*; **c, d** *Labrisomus nuchipinnis*; **e, f** and *Bathygobius soporator*; **g, h** Ag-NOR-bearing chromosome pairs are highlighted.

chromosomes of the karyotype was far less pronounced. Ribosomal sites were on the terminal portions of the short arms on chromosome pair 4. C-banding showed discrete heterochromatic regions in the centromeric regions of most chromosomes and telomeric regions of some acrocentric pairs (Fig. 1g, h).

Discussion

Though many perciform families display a conserved karyotype pattern, with $2n=48$ acrocentric chromosomes, some groups demonstrate dynamic tendencies in relation to chromosome evolution (Molina 2007). Much of identifiable chromosome diversity is attributed to pericentric inversions, the most common mechanism of chromosome evolution in this order (Galetti et al. 2000, 2006).

Representatives of the suborder Blennioidei (e.g., Carbone et al. 1987) and Gobioidaei (e.g., Arai and Sawada 1974, 1975; Thode et al. 1988; Oliveira and Almeida-Toledo 2006) stand out for their greater karyotype variability and diversity. This includes species with conserved karyotypes and those that are highly diversified.

Within the Blennioidei, the Blenniidae, a monophyletic family, is divided into six tribes including Salariini and Parablenniini which, in turn, include the Atlantic species *O. trinitatis* and *S. cristata* respectively (Nelson 2006). Comparisons of mitochondrial DNA sequences in samples of *Ophioblennius* Gill, 1860 collected throughout the Atlantic suggest that the genus consists of six distinct lineages. One of these corresponds to species found in the Pacific, while the rest are recorded in the biogeographic provinces of the Atlantic: Brazilian, Caribbean, Mid-Atlantic, Sao Tome and Azores/Cape Verde (Muss et al. 2001). Chromosome characteristics reported here for *O. trinitatis* are the first for the genus, exhibiting $2n=46$, $6m+12st+28a$ and $FN=64$. The relatively low diploid number and higher fundamental number in relation to the mean of other species of Blenniidae (Table 1), as well as the presence of large metacentric chromosomes, suggests pericentric inversion events and the occurrence of Robertsonian translocation involving two of its chromosome pairs. In turn, *S. cristata*, while also belonging to the family Blenniidae, has a distinct karyotype of $2n=48$, $4st+44a$ and $FN=52$. Thus, *S. cristata* differs from *O. trinitatis* in that it contains an extra pair of chromosomes, lacks metacentric chromosomes and has different numbers of subtelocentric and acrocentric chromosomes in the karyotype. The karyotype of the *S. cristata* population studied here differs from the karyotypes previously described for the coastal population of Rio de Janeiro (SE Brazil), with $2sm+46st/a$ (Brum et al. 1994), and the Mediterranean population, with $2st+46a$ (Vitturi et al. 1986). Nevertheless, despite the growing number of discordant karyotype descriptions between populations on the NE and SE coasts of Brazil, one cannot rule out that these differences may arise from the difficulty in precisely defining types of cryptic chromosomes in the karyotype of this species.

In spite of displaying relative diversity in chromosome structure, only 18.5% of Blennioidei species exhibit differences in the basal diploid number, $2n=48$ chromosomes. As shown in table 1, diploid numbers for representatives of this suborder vary

between $2n=40$, found in *Dasson trossulus* (Jordan & Snyder, 1902) (Arai and Shiotsuki 1974) and $2n=52$ in *Gobius niger* Linnaeus, 1758 (Vitturi and Catalano 1989), but with a conspicuous modal value of $2n=48$.

In contrast to Blennioidei, suborder Gobioidae shows much more dynamic karyotype evolution, demonstrating highly variable karyotype patterns, where the diploid number ranges from $2n=30$ for *Neogobius eurycephalus* (Kessler, 1874) (Ene 2003), to $2n=62$ in *Mogurnda mogurnda* (Richardson, 1844) (Nogusa 1960). Cytogenetic data for 95 species show that only 9.6% have $2n=48$ chromosomes, whereas the highest frequencies observed correspond to $2n=46$ in 40% of species investigated, and $2n=44$ in 32% (Table 1). As such, both Gobioidae species studied here are included in the group showing $2n=48$ chromosomes, *L. nuchipinnis* with $2st+46a$ and $FN=50$ and *B. saporator* with $2m+6st+40a$ and $FN=56$. Thus, *B. saporator* differs from *L. nuchipinnis* in the presence of metacentric chromosomes and different numbers of subtelocentric and acrocentric chromosomes in the karyotype.

Among chromosome rearrangements involved in karyotypic differentiation of Gobiidae, Robertsonian fusions stand out, and are likely the most common event in this group (Amores et al. 1990; Galetti et al. 2000). However, other more complex changes in karyotypic structure (Thode et al. 1988; Vitturi and Catalano 1989; Caputo et al. 1997; Caputo et al. 1999), as well as the presence of different sex chromosomes (e.g., Pezold 1984; Baroiller et al. 1999), can also be observed, corroborating the high dynamic evolution that characterizes suborder Gobioidae. It has been suggested that the baseline/ancestral karyotype for Gobiidae would consist of $2n=46$ acrocentric chromosomes (Vasil'ev and Grigoryan 1993), from which an increase in bi-brachial chromosomes would characterize more derived karyotypes. Based on this proposal, *B. saporator* ($FN=56$) would experience a greater number of structural rearrangements during its karyotypic evolution process in relation to *L. nuchipinnis* ($FN=50$).

Location and frequency of Ag-NOR sites are efficient cytotaxonomic markers in many groups of fish (Caputo 1998). Among species of Gobiidae, at least six different arrangement patterns for nucleolar organizer regions have been identified (Fig. 2), which supports the occurrence of intense karyotypic diversification mechanisms in this group. Thus, Ag-NOR sites can be found (a) in the telomeric region on the short arm of a single pair of acrocentric chromosomes, as in *Gobius fallax* Sarato, 1889 (Thode et al. 1983) and *Gobius paganellus* Linnaeus, 1758 (Caputo 1998); (b) in the telomeric region on the long arm of a single pair of acrocentrics, such as in *Zosterisessor ophiocephalus* (Pallas, 1814) (Caputo 1998); (c) in the interstitial/pericentromeric region on the long arm of a single pair of acrocentric chromosomes, as seen in *Proterorhinus marmoratus* (Pallas, 1814) (Ráb 1985) and *Gobius cobitis* Pallas, 1814 (Caputo 1998); (d) in the telomeric region on the short arm of a single subtelocentric pair, described in *B. saporator*; (e) in the interstitial/pericentromeric region on the long arm of a single metacentric pair, observed in *N. eurycephalus* (Ene 2003); and (f) in the telomeric regions on the short arms of two acrocentric chromosome pairs, recorded in *Gobiusculus flavescens* (Fabricius, 1779) (Klinkhardt 1992).

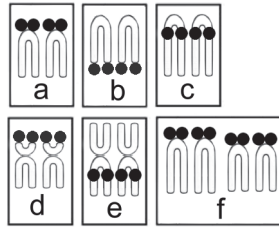


Figure 2. Ag-NOR phenotypes **a–f** described in species of Gobiidae. Ag-NORs sites described in the karyotypes of Gobiidae species were found **a** in the telomeric region on the short arm of a single pair of acrocentric chromosomes **b** in the telomeric region on the long arm of a single pair of acrocentrics **c** in the interstitial/pericentromeric region on the long arm of a single pair of acrocentric chromosomes **d** in the telomeric region on the short arm of a single submetacentric pair **e** in the interstitial/pericentromeric region on the long arm of a single metacentric pair and **f** in the telomeric regions on the short arms of two acrocentric chromosome pairs.

Few data are available on ribosomal sites for Labrisomidae. Ag-NORs in *L. nuchipinnis* exhibit the phenotype (b) described above, in addition to both species of Blennioidei, *O. trinitatis* and *S. cristata*, which may suggest an ancestral condition for this location.

In contrast, other chromosome characteristics, such as C-positive heterochromatin distribution, may be more conserved. This occurs in several species of Perciformes where discrete blocks are preferentially located in the centromeric/pericentromeric regions of chromosomes (Molina 2007). This pattern is repeated in *S. cristata*, *O. trinitatis* and *L. nuchipinnis*, as well as in some Gobiidae, such as *G. cobitis*, *Z. ophioccephalus* and *N. eurycephalus* (e.g. Caputo et al. 1997; Ene 2003). In *B. soporator*, in addition to centromeric/pericentromeric regions, heterochromatic sites are also observed in terminal regions of some chromosomes. This arrangement has already been described for other Gobiidae, including *G. paganellus* and *G. niger*, where pericentromeric and telomeric heterochromatic regions are distributed among almost all chromosomes (Amores et al. 1990; Caputo et al. 1997).

Moreover, karyotypic diversity present in Gobioidae is increased by the occurrence of chromosome polymorphisms frequently observed in this group. This is particularly evident in several examples of intraspecific karyotypic variability, as well as polymorphisms involving different types of chromosome rearrangements, such as in *G. niger* (Vitturi and Catalano 1989; Caputo et al. 1997) and *G. fallax* (Thode et al. 1988). Data obtained for the paedomorphic Gobiidae *Aphia minuta* (Risso, 1810) also show variations in the diploid number and chromosome formula, resulting in five different cytotypes ($2n=41-44$ and $FN=42-44$) (Caputo et al. 1999). Similar karyotypic variability was reported in *N. eurycephalus*, where three specific cytotypes ($2n=30, 31$ and 32) were associated to the occurrence of centric fusions (Ene 2003). All these examples demonstrate clear chromosomal dynamism, with possible transitions to new karyotype patterns.

In fact, karyotypic diversity among Blennioidei and Gobioidae seems to accompany phyletic diversification of these groups. This is a result of vicariant factors (Pampoulie et al. 2004) and could be favored by their low dispersive potential (Fanta 1997), as well as ecological specificities that favor population fractionation in this family (Huyse et al. 2004). The present study also highlights the importance of ribosomal sites as effective chromosomal markers in the further cytogenetic studies in gobioid species.

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