Chromosomal studies of three vulnerable marine fishes from west coast of India

Basdeo Kushwaha, Ravindra Kumar*, N S Nagpure, Satish K Srivastava, V S Basheer, M K Anil¹ & W S Lakra Molecular Biology and Biotechnology Division, National Bureau of Fish Genetic Resources, Canal Ring Road, P.O. Dilkusha, Lucknow- 226 002, India.

¹Senior Scientist, Vizhinjam Research Centre of CMFRI, P.B. No. 9 Vizhinjam P.O., Thiruvananthapuram-692 521, India

*[E-mail:Ravindra.Scientist@gmail.com,ravindrakumar@scientist.com, rkumar@nbfgr.res.in]

Received 21 August 2009; revised 15 January 2010

Cytogenetic profiling was carried out in three vulnerable marine ornamental fishes, namely *Thalassoma lunare*, *Zanclus cornutus* and *Arius subrostratus*, using silver nitrate, chromoycin A3 CMA3 staining and C- banding techniques to study the variation in localization of NORs and C-bands. Karyotype analyses of these species revealed a diploid chromosome number of 48, all acrocentrics, in *T. lunare* and *Z. cornutus*. In *A. subrostratus*, however, the diploid chromosome number was found to be 58 consisting of 22 metacentric, 16 submetacentric, 10 subtelocentric and 10 telocentric chromosomes with fundamental arm number of 96. The silver stained NORs were observed on 3 pairs of chromosomes in *T. lunare*, whereas other two species possessed NOR on single pair of chromosome. Within the species, there was complete concordance in number and position of NORs as detected by AgNO3 and CMA3 staining. Prominent constitutive heterochromatic bands were detected on 4, 2 and 3 pairs of chromosomes, respectively, in *T. lunare*, *Z. cornutus* and *A. subrostratus*. There was variation found in the number and position on NORs and C-bands among these species, which could be used as species-specific markers. This study describes for the first time the cytogenetic profiling in *Z. cornutus* and *A. subrostratus*.

[**Keywords:** Ag-NORs; C-bands; Chromomycin A₃; Cytogenetics; Marine fishes].

Introduction

Though India has rich diversity of ichthyofauna, merely about 40 species from marine ecosystems have been cytogenetically characterized¹. In the present scenario of WTO and IPR regime, genetic characterization of fish fauna has gained utmost importance. The karyotypic information can throw light on genetic characterization, phylogenetic relationships among species and karyotype evolution in fishes. In view of the above, the present cytogenetic studies have been carried out in three marine fish species viz., *Thalassoma lunare, Zanclus cornutus* and *Arius subrostratus* so as to facilitate conservation and management of natural stocks of these species.

Materials and Methods

Six live specimens each of T. lunare and

*Corresponding author:

Dr. Ravindra Kumar, National Bureau of Fish Genetic Resources Canal Ring Road, P.O. Dilkusha, Lucknow - 226 002, India Phone +91-522-2442440, 2442441, Fax: +91-522-2442403, E-mail: Ravindra.Scientist@gmail.com, ravindrakumar@scientist.com, rkumar@nbfgr.res.in

Z. cornutus were collected from Kovalam area, Kerala whereas specimens of A. subrostratus (n=5) were collected from Vypeen Island, Cochin, Kerala. The average wet weight of the specimens of T. lunare was 92 g (range 75-100) whereas those of Z. cornutus and A. subrostratus were 21.8 g (range 20-25) and 23 g (range 20-25), respectively. All the specimens were in juvenile stage and the sex was unidentifiable by visual examination.

For preparing chromosome slides, the specimens were administered intramuscularly with 0.05% colchicine @1.0 ml/100 g body weight and were sacrificed after 2 h for dissecting out the kidney tissue. The tissue was homogenized in 0.56% KCl and further processed for chromosome preparations using fixation (acetic acid-methanol fixation) and flame drying technique. The chromosomes were stained with 4% Giemsa in Phosphate buffer (pH 6.8). Approximately, 200 chromosome complements analyzed for establishing the were modal chromosome number and characteristic chromosome morphology of each species. For karyotyping, chromosomes were grouped into metacentric (m),

submetacentric (sm), subtelocentric (st) and telocentric (t) as per the classification proposed by Levan *et al.*²

Localization of nucleolar organizer regions (NORs) was studied by both silver nitrate (AgNO₃) impregnation and chromoycin A3 CMA₃ staining techniques. Ag-NOR staining was performed according to the method of Howell and Black³. For CMA₃ staining, the method described by Ueda et al.⁴ was followed. C-banding was carried out as per the technique described by Sumner⁵ with staining of slides by propidium iodide (5µg/ ml in DDW) instead Giemsa as described bv et al.6 A particular band pattern was determined by studying a minimum of 20 metaphase spreads per specimen.

Results

Chromosome number was conserved in both the perciform species, viz. *T. lunare* and *Z. cornutus*, as evidenced by the model diploid number of 48 acrocentric chromosomes (FN= 48) (Figs. 1a, b). In the catfish *A. subrostratus*, however, the diploid chromosome number was 58 with karyotype formula 22m + 16sm + 10st + 10t (FN= 96) (Fig. 1c).

The silver-stained NOR indicates the transcriptional activity of NORs. The Ag-NOR staining of chromosomes revealed variation in number and position of NORs among these species.

In *T. lunare*, the silver stained NORs were present on terminal portions of 3 pairs of chromosomes, i.e. p arm of 4th, 12th and 18th chromosomes (Fig. 2a). In both *Z. cornutus* and *A. subrostratus*, the NORs were present on single pair of chromosome (Figs. 2b, c) and the position was terminal on 10th chromosome in the former and on 5th submetacentric pair in the later.

The CMA₃ staining of metaphase complements in these marine species showed bright fluorescent signals at the same position (Figs. 2a-c), which were positive for AgNO₃ staining, thus, confirmed the position of active transcribing zones of NOR-bearing chromosomes. In *Z. cornutus*, a prominent CMA₃ band present on pericentric region of a chromosome showed likely association with C-band.

The C-banding technique revealed presence of constitutive heterochromatin (CH) blocks on few chromosomes in these species and the pattern was found to be species-specific. In *T. lunare*, the C-bands were present on 4 pair of chromosomes at centromeric positions (Fig. 2a), while in *Z. cornutus*, C-bands were located at centromeric position on one pair of chromosomes (Fig. 2b). In *A. subrostratus*, prominent C-bands were present at the p arm of the three pair of chromosomes (Fig. 2c). The variation in positions of heterochromatic blocks could be used as species-specific marker for identification of these species.

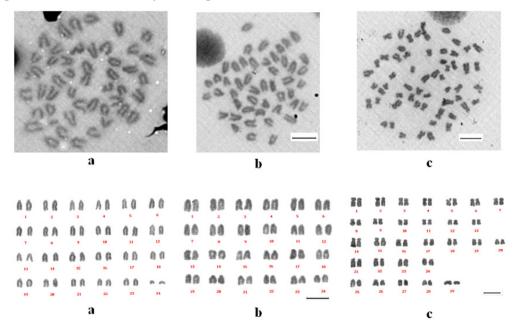


Fig. 1—Giemsa stained metaphase spreads and karyotypes of: (a) T. lunare, (b) Z. cornutus and (c) A. subrostratus. Bar size: 5 µm.

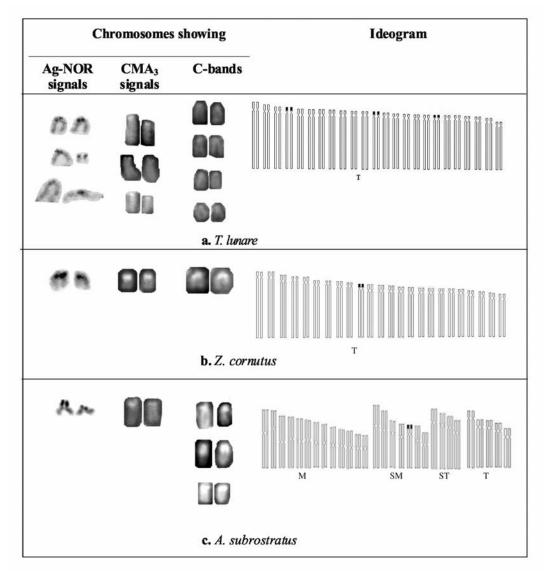


Fig. 2—Chromosomes showing silver and CMA₃ stained NORs, C-bands and ideogramatic representation in: (a) *T. lunare*, (b) *Z. cornutus* and (c) *S. subrostratus*. Dark regions in ideogram indicate the NOR-bearing chromosomes

Discussion

The family Labridae (Wrasses) contains approximately 500 species distributed into 60 genera, and so far only 47 species (belonging to 21 genera) of them are cytogenetically studied⁷⁻¹². Under genus species Thalassoma, around 62 are listed (www.fishbase.org, version 4/2009) and only 6 of them were karyotyped. In all the earlier reports on T. lunare, the diploid chromosome number was found to be 48^{11,13}.

To date, no cytogenetic investigation has been undertaken in *Z. cornutus*. The species also possessed 2n= 48, with all acrocentric chromosome, thus, carrying the characteristics of the higher group. So

far, chromosomal studies are available only for 9 species (7 genera) of Ariidae family^{7-9,14-17}.

Evolutionary stasis has often been referred to as a common phenomenon, whether at a higher or a lower degree, in the karyotypic evolution of Perciform species¹⁸. In family Labridae, the degree of karyotypic differentiation (number of changes in FN values) is inversely related to the dispersive potential provided by the extent of pelagic larval duration of each species¹⁸⁻²⁰.

In the family Labridae, although most species have been reported with 2n=48, the number of chromosome arms (FN) is often higher (48–90), indicating a predominant occurrence of pericentric inversions within this group²¹. The representative fish species of the family Ariidae show a karyotypic pattern of 2n=54–58 and high chromosome arm numbers, FN>80^{22,23}, which is very similar to the most probable ancient karyotype of the Siluriformes²⁴.

The presence of single NOR pair in fishes, in general, was considered to be plesiomorphic or primitive condition and multiple NORs as apomorphic or derived condition²⁵. The remarkable karyotypic conservativeness detected in the present study is similar to that observed in other perciform species previously studied, regarding both the number of acrocentric chromosomes and NOR location.

The GC-rich active regions of NORs are revealed by CMA₃ staining technique and a positive correlation was observed between AgNO₃ and CMA₃ staining in these undertaken marine species that indicated presence of active NOR in these species.

Positive C-bands identify regions heterochromatins, which consist of repetitive DNA sequences and are not transcriptionally active²⁶. No earlier report on the C-banding pattern in these undertaken marine species is available. In the present study, high genetic diversity with respect to the heterochromatic bands was observed in the genome of three marine fishes. In T. lunare, the bands were located on 4 pair of chromosomes at their centromeric positions, whereas the other perciforme fish Z. cornutus possessed prominent C-band on only one pair of chromosome at centromeric position. The centromeric positions of C-bands have been reported in large number of fish species^{27,28}. Large variations in number and localization of C-bands have also been reported in our study on six freshwater fish species belonging to the genus *Tor* (unpublished results).

The distribution of C-bands on the different chromosomes may be due to the pericentric inversion²⁹. C-band positive segments in the terminal and centromeric positions have been observed in flatfishes³⁰ and in Labeo rohita and Cirrhinus mrigala³¹. Almeida-Toledo et al.³² and Uwa³³ observed such type of observations in other fish species and suggested that the large C-banded heterochromatic region could have been formed by tandem duplication of heterochromatic DNA that may considered be species specific marker chromosome. Similar to our finding, heterochromatin

blocks were also detected at the centromeric position on most of the chromosome pairs in Brazilian marine fishes belonging to the Sciaenidae and Sparidae groups³⁴. Positive correlation between C-bands and NOR bearing chromosomes were observed in all species under study which seems to be a common feature in other fish species³⁵⁻³⁷.

The distribution of C-banded heterochromatin allows for a more accurate pairing of homologous chromosome of the metacentric, submetacentric and subtelocentric series in *A. subrostratus*. Another characteristic about the observed C-bands were their localization at centromeric regions in both the perciformes species and at telomeric position in *A. subrostratus*. This indicated that the heterochromatic regions in *A. subrostratus* might have evolved due to breaking and rejoining of chromosomes during karyoevolution.

Detailed information related to the differences in chromosome morphology and the position as well as number of NORs and C-bands in the undertaken species suggested that these cytotaxonomic markers could be utilized in identification and characterization of genetic variability between the species and their hybrids. Moreover, the karyological data supplemented with technological progress in molecular cytogenetics will offer important evidences for the classification of these fish species.

Acknowledgements

Authors are thankful to the Director, NBFGR, Lucknow and Scientist in-charge Vizhinjam Research Center of CMFRI for providing facilities to carry out this part of work.

References

- Nagpure, N.S., Kumar, R., Srivastava, S.K., Kushwaha, B., Gopalakrishnan, A. & Basheer, V.S., Cytogenetic characterization of two marine ornamental fishes, *Chaetodon collare* and *Stegastes insularis*, *J. Mar. Biol. Ass. India*, 48 (2006) 267-269.
- 2 Levan, A., Fredga, K. & Sandberg, A. A., Nomenclature for centromeric position on chromosomes. *Hereditas*, 52 (1964) 201–220.
- 3 Howell, W.M. & Black, D.A., Controlled silver-staining of nucleolus organizer regions with a protective colloidal developer: a 1-step method, *Experientia*, 36 (1980) 1014.
- 4 Ueda, T., Irie, S. & Kato, Y., Longitudinal differentiation of metaphase chromosomes of Indian muntjac as studies by restriction enzyme digestion, in situ hybridization with cloned DNA probes and distamycin A plus DAPI fluorescence staining, Chromosoma (Berl.), 95 (1987) 251– 257.

- 5 Sumner, A.T., *Chromosome Banding*. Cambridge: Unwin Hyman Inc, 434 p (1990).
- 6 Fontana, F., Tagliavini, J., Congiu, L., Lanfredi, M., Chicca, M., Laurenti, C. & Rossi, R., Karyotypic characterization of great sturgeon, *Huso huso*, by multiple staining techniques and fluorescent *in situ* hybridization, *Mar. Biol.*, 132 (1998) 495–501.
- 7 Arkhipchuk, V.V., Chromosome database. *Database of Dr. Victor Arkhipchuk*, (1999).
- 8 Brum, M.J.I., Cytogenetic studies of Brazilian marine fish, *Braz. J. Genet.*, 19(3) (1996) 421-427.
- 9 Klinkhardt, M.B, Tesche, M. & Greven, H., Database of fish chromosomes, Westarp Wissenschaften, Magdeburg (1995).
- 10 Manna, G.K., Fish cytogenetics related to taxonomy, evolution and monitoring aquatic genotoxic agents. p. 21-46 (1989). In P. Das and A.G. Jingran (eds.) Fish Genetics in India. Proceedings of the Symposium on Conservation and Management of Fish Genetic Resources of India. 11-13 April, 1986.
- 11 Ojima, Y. & Kashiwagi, E., Further Studies of the Chromosomes of the Labridae (Pisces), *Proc. Japan Acad.*, 56 Ser. B (1989) 328-331.
- 12 Vasil'ev, V.P., Chromosome numbers in fish-like vertebrates and fish, *J. Ichthyol.*, 20(3) (1980) 1-38.
- 13 Ojima, Y. & Kashiwagi, E., A karyotype study of eleven species of labrid fishes from Japan, *Proc. Japan Acad.*, 55B (1979) 280-285.
- 14 Hinegardner, R., & Rosen, D.E., Cellular DNA content and the evolution of teleostean fishes, *Am. Nat.*, 106(951) (1972) 621-644.
- 15 Gomes, V., Ngan, P.V., M.J. de A. & Passos, C.R., The karyotype of a marine catfish, *Bagre bagre*, from Brazil, *Jap. J. Ichthyol.*, 37(3) (1990) 321-323.
- 16 Leggatt, R.A. & Iwama, G.K., Occurrence of polyploidy in the fishes, Rev. Fish Biol. Fish., 13 (2003) 237-246.
- 17 NBFGR, Fish chromosome atlas, National Bureau of Fish Genetic Resources Special Publication, No. 1. Lucknow, India, (1998) 332 p.
- 18 Molina, W.F., Chromosomal changes and stasis in marine fish groups, In: Fish cytogenetics (Pisano E, Ozouf-Costaz C, Forest F and Kapoor BG, eds.). Science Publishers, Enfield (2006) 69-110.
- 19 Molina, W.F. & Galetti, P.M.Jr., Multiple pericentric inversions and chromosomal divergence in the reef fishes Stegastes (Perciformes, Pomacentridae), *Genet. Mol. Biol.*, 27 (2004) 543-548.
- 20 Sena, D.C. & Molina, W.F., Robertsonian rearrangements and pericentric inversions in Scaridae fish (Perciformes), *Genet. Mol. Res.*, 6 (2007) 575-580.
- 21 Alvarez, M.C., Garcia, E. & Thode, G., Contribution to the karyoevolutive study of the Labridae (Perciformes). The karyotypes of *Ctenolabrus rupestris* and *Symphodus* ocellatus, Caryologia, 39(1986) 353–357.
- 22 Rishi, K.K. & Singh, J., Karyological studies on two Indian estuarine catfishes, *Plotosus canius* Ham. and *Pseudeutropius atherinoides* (Bloch), *Caryologia*, 36 (1983) 139–144.

- 23 Gomes, V., Phan, V. N. & Passo, M.J.A.C., Karyotypes of three species of marine catfishes from Brazil, *Bolm. Inst.* oceanogr. USP, 42 (1994) 55–61.
- 24 LeGrande, W.H., Chromosomal evolution in North American catfishes (Siluriformes: Ictaluridae) with particular emphasis on the Madtoms, *Noturus*, *Copeia*, 1981 (1) (1981) 33–52.
- 25 Gold, J.R. & Amemiya, C.T., Cytogenetic studies in North American minnows (cyprinidae) XII. Pattern of chromosomal nucleolus organizer region variation among 14 species, *Canadian J. Zool.*, 65 (1986) 1869–1877.
- 26 Sumner, A.T., The mitotic chromosome. In: *Advances in Genome Biology*. Jai Press Inc., 5 (1998) 211-261.
- 27 Gold, J.R., Li, Y.C., Shipley, N.S. & Powers, P.K., Improved methods for working with fish chromosomes with a review of metaphase chromosome banding, *J. Fish Biol.*, 37 (1990) 563-575.
- 28 Rishi, K.K. & Mandhan, R.P., Analysis of C-banded heterochromatin in chromosomes of *Labeo rohita* (Ham) (Cyprinidae), *Chromosome Information Service*, 48 (1990) 3-4
- 29 Sen, S. & Sharma, T., Role of constitutive heterochromatin in evolutionary divergent: results of chromosome banding and condensation in inhibition studies in *Mus musculus*, *Mus booduga* and *Mus dunni*, *Evol.*, 37(3) (1983) 628-636.
- 30 Azevedo, M.F.C, Oliveria, C., Pardo, B.G., Martinez, P. & Foresti, F. Cytogenetic characterization of six species of flatfishes with comments to karyotype differentiation patterns in Pleuronectiformes (Teleostei), J. Fish Biol., 70 (2007) 1-15.
- 31 Khuda-Bukhsh, A.R. & Chakrabarty, C., Localization of C-band heterochromatin in metaphase chromosomes of two species in Indian major carp, *Inland Fish Soc. India*, 26 (1994) 44-46.
- 32 Almeida-Toledo, L.F., Foresti, F. & Toledo, S.A., Constitutive heterochromatin and nucleolus organizer regions in the knifefish *Apteronotus albifrons* (Pisces, pteronotidae), *Experientia*, 37 (1981) 953-954.
- 33 Uwa, H., Karyotype evolution and geographical distribution in the ricefish, genus *Oryzias* (Oryziidae), In: T. Uyeno, R. Arai, T. Taniuchi, and K. Matsuura (eds), *Indo Pacific Fish Biology*, Tokyo, Ichthyol. Soc. of Japan, p. 867-876 (1986).
- 34 Accioly, I.V. & Molina, W.F., Cytogenetic studies in Brazilian marine Sciaenidae and Sparidae fishes (Perciformes), *Genet Mol Res.*, 7(2) (2008) 358-70.
- 35 Pendas, A.M, Moran, P. & Garcia-Vázquez, E., Ribosomal RNA genes are interspersed throughout a heterochromatic chromosome arm in Atlantic salmon, *Cytogenet. Cell Genet.*, 63 (1993) 128-130.
- 36 Martinez, J.L., Moran, P., Garcia-Vazquez, E. & Pendas, A.M., Chromosomal localization of the major and 5S rRNA genes in the European eel (*Anguilla anguilla*), *Cytogenet. Cell Genet.*, 73 (1996) 149-152.
- 37 Wasko, A.P. & Galetti, P.M. Jr., Mapping 18S ribosomal genes in fish of the genus *Brycon* (Characidae) by fluorescence *in situ* hybridization (FISH), *Genet. Mol. Biol.*, 23(1) (2000) 135-138.