

Original Research Article

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Chromosomal and Banding Studies in *Sphaerium indicum* (Bivalvia: Mollusca)

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ABSTRACT

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During the present study, mitotic chromosomes of freshwater snail *Sphaerium indicum* were studied using gills. The observed diploid chromosome number was $2n=30$ consisting of 26 metacentric pairs and 04 submetacentric pairs respectively. Chromosomes were studied using Giemsa staining and two banding techniques viz. C-banding and Ag-NOR banding. Preliminary C-band studies revealed the presence of terminal blocks besides the presence of centromeric heterochromatin. The nucleolar organizer region (NOR) was found on one chromosome pair i.e. on the submetacentric chromosome pair no. 5. The cytogenetic relationship between *S. indicum* and other species of Veneroida was also discussed.

Introduction

Molluscs are abundant, varied and widespread. This and the diversity of their karyotypes make them particularly suited to cytogenetic research. Cytogenetics studies encompass different levels of biological organization ranging from the morphological to the molecular, depending on the applicable technology. Chromosomes can be studied as a morphological manifestation of the genome in terms of their microscopically visible size, shape, number and behavior during meiosis and mitosis. At another level, banding studies reveal finer details of chromosomal morphology. Studies of natural or induced

deviation from the diploid chromosome number also contribute significant cytogenetic data. Among bivalves, the first investigation concerned the chromosome number (Patterson, 1969). Then, pre-treatment with colchicine and hypotonic treatment, combined with the air drying technique, permitted accurate assessments of chromosome morphology (Nakamura, 1985). Later, the development of banding techniques which allowed chromosome identification in karyotypes began to be applied in bivalves (Thiriot-Quievreux, 1994).

S. indicum is an abundant mollusc species in the Himalayan streams and riverlets within the

elevation range of 800-1600m. The preferred habitat is sandy gravel or muddy substrate closed to the banks and slow running water bodies. It belongs to Phylum –Mollusca, Class-Bivalvia, Order- Veneroida, Family - Pisiidae and Genus – *Sphaerium* species *indicum*.

Taxonomic characters includes: Shell in equilateral, right valve with a single well developed cardinal, Mantle very thin and without any papillae on the edge. It is commonly known as Fingernail Clam.

Materials and Methods

The snails were collected from slow running fresh water streams. Snails were taken alive to the laboratory, then maintained in tanks of aerated water and fed continuously to promote growth. Chromosome preparations were obtained from the gills tissue.

After taxonomic verification of each snail, were floated for 24 hrs in 0.1% colchicine. Snails were dissected and their gills removed and treated with 0.07% KCl as hypotonic solution, at room temperature for 30 min. The tissues after hypotonic treatment were fixed in Carnoy's fixative. The slides were stained with 4% Giemsa buffer solution for 30 min. scanning and photomicrography of the slides was done using Nikon YS100 binocular research microscope and Samsung SDC-313 camera respectively. Well spread suitable mitotic stages were photomicrographed at a magnification of 1000x. The morphological classification of chromosomes proposed by Levan *et al.*, (1964) has been followed to categorize the chromosomes as metacentrics, submetacentrics, subtelocentrics or telocentric. The chromosomes were classified into uniarmed and biarmed following Chen and Ebelling (1971) to calculate the fundamental arm number (FN). Idiogram and Histogram was prepared.

Observations

A total of 40-50 metaphase stages were selected to establish the diploid chromosome number. The basic chromosome number was found to be $2n=30$ (Fig. 1). All chromosomes in the karyotypes were biarmed and of two types, that is, metacentric and submetacentric type (Fig. 2) Haploid formula for the complement was calculated as $n=13M+2SM$ and the corresponding fundamental arm number was calculated as $NF=60$ (Table 1).

The mean total haploid length and total diploid length was 12.73 μm and 25.46 μm respectively. Histogram and Idiogram was also prepared (Fig. 3 and 4) In C- Banding, Chromosome pair numbers 1, 2, 3, 4, 5, 6, 10, 11 and 15 showed telomeric C-bands, while the chromosome pair number 7, 8, 9, 12, 13 and 14 showed centromeric C-bands (Fig. 5 and 6) while in NOR- Banding, one pair of darkly stained NOR was observed in a short arm of submetacentric chromosome pair (5th pair) (Fig. 7 and 8).

Results and Discussion

Although Bivalves are an important component of the marine and freshwater fauna, their chromosomal studies have received a little consideration than they actually deserve. Karyotypic details have been documented in several animal groups including a few Bivalvia species (Cornet and Soulard, 1990; Wada and Komaru, 1993; Insua *et al.*, 1998; Martinez-Lage *et al.*, 2002; Lopez-Pinon *et al.*, 2005; Insua *et al.*, 2006). Karyotypes with a majority of metacentric-submetacentric chromosomes are also characteristic of most bivalve species (Thiriot-Quievreux, 2002), but this feature is not a general rule among invertebrates (White, 1973). Out of 10,000 described species of Bivalvia only 139 species has been studied cytogenetically till date.

Fig.1 Metaphase complement from gill

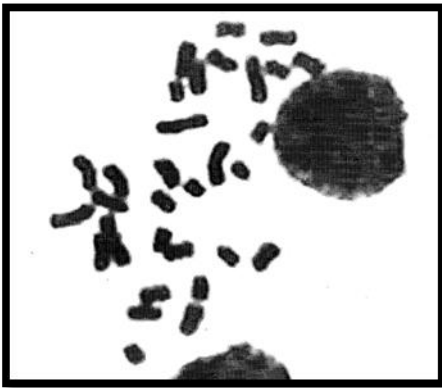


Fig.2 karyotype showing $2n=30$ (26m+4sm)
Tissue of *S. indicum* ($2n=30$)

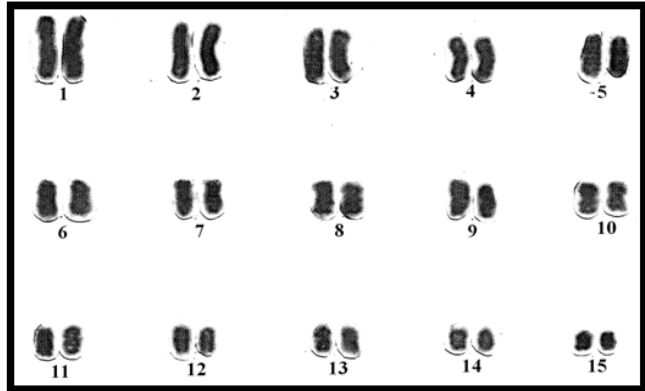


Fig.3 Histogram

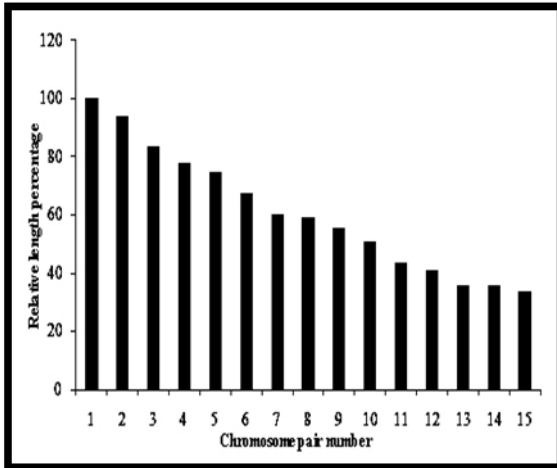


Fig.4 Idiogram

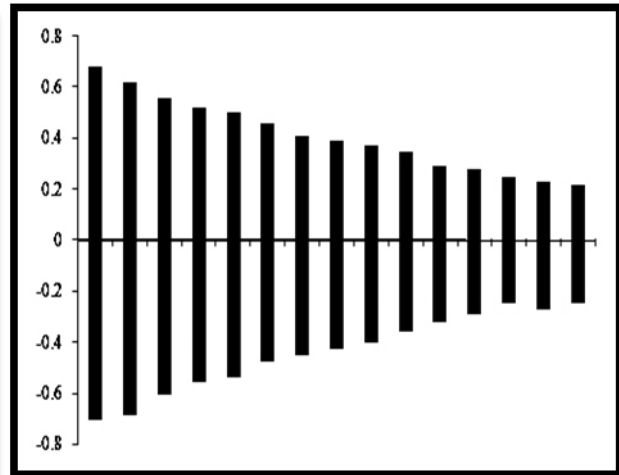


Fig.5 C-banded metaphase complement



Fig.6 Karyotype of C-banded metaphase complement

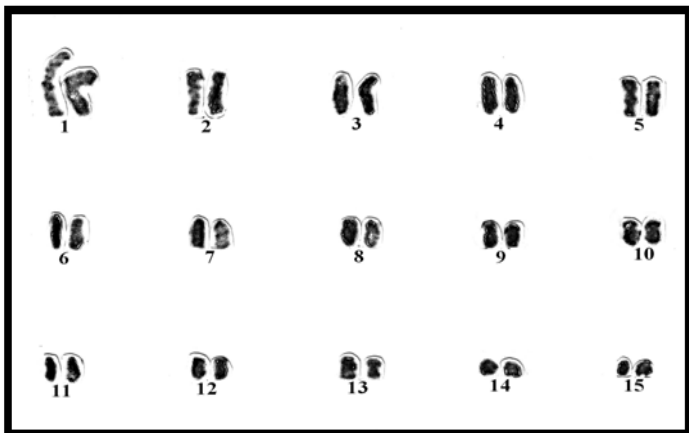


Fig.7 NOR banded metaphase complement

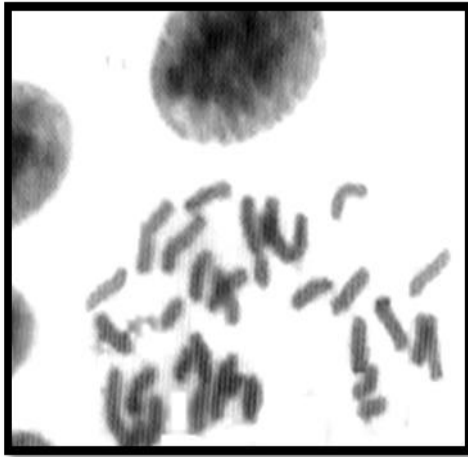


Fig.8 Karyotype of NOR-banded metaphase Complement

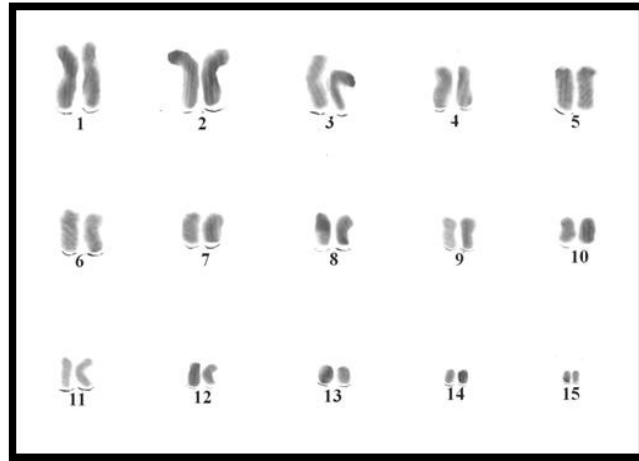


Table.1 Morphometric data of karyotype of *S. indicum* showing $2n=30$ (26m+4sm)

Chromosome pair no.	Mean length of the short arm (p) in μm	Mean length of the long arm (q) in μm	Absolute length (p+q) of the chromosome in μm	Arm ratio (q/p)	Relative length percentage	Total complement length percentage	Centromeric index	Nomenclature
1	0.68	0.71	1.39	1.04	100.00	5.46	48.92	Metacentric
2	0.62	0.69	1.31	1.11	94.24	5.15	47.32	Metacentric
3	0.56	0.60	1.16	1.07	83.45	4.56	48.27	Metacentric
4	0.52	0.56	1.08	1.07	77.69	4.24	48.14	Metacentric
5	0.50	0.54	1.04	1.08	74.82	4.08	48.07	Submetacentric
6	0.46	0.48	0.94	1.04	67.62	3.69	48.93	Metacentric
7	0.41	0.45	0.86	1.09	60.43	3.38	47.67	Metacentric
8	0.39	0.43	0.82	1.10	58.99	3.22	47.56	Submetacentric
9	0.37	0.40	0.77	1.08	55.39	3.02	48.05	Metacentric
10	0.35	0.36	0.71	1.02	51.07	2.79	49.29	Metacentric
11	0.29	0.32	0.61	1.10	43.88	2.39	47.54	Metacentric
12	0.28	0.29	0.57	1.03	41.00	2.24	49.12	Metacentric
13	0.25	0.25	0.50	1.25	35.97	1.96	50.00	Metacentric
14	0.23	0.27	0.50	1.17	35.97	1.96	46.00	Metacentric
15	0.22	0.25	0.47	1.13	33.81	1.85	46.80	Metacentric

The diploid chromosome number ranges between 12-62, with 28 being the most common (28.8%); followed by 38 (25.2%)

and 20 (17.3%). Cytogenetic studies conducted on bivalves have revealed significant variation in the number of

chromosomes. In a review (Nakamura, 1985), it was reported that the order Veneroida, has diploid chromosome number from 24 to 48 and the most common chromosome number was $2n=38$. However, deviations from this pattern appeared within individual families.

In family Pisidiidae (Bivalvia: Veneroida), three species has been studied cytologically till date, viz. *Sphaerium corneum*, $2n=30$ with 26m+4sm chromosomes (FN=60) (Petkeviciute *et al.*, 2006; Korinkova and Kral, 2011); *S. nucleus*, $2n=30$ with 28m+2sm chromosomes (FN=60) (Korinkova and Kral, 2011) and *S. nitidum*, $2n=44$ with 24m + 8sm + 12st chromosomes (FN=88) (Barsiene and Barsyte, 2000). In present study for the first time, chromosomal and banding studies of *S. indicum* has been done. The diploid complement of $2n=30$ with 26m+4sm chromosomes (FN=60) was observed in *S. indicum*, showing the presence of only biarmed chromosomes. This may be due to Robertsonian translocation (centric fission) of chromosomes or polyploidy as there was almost doubling in the chromosome as compared to other species.

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