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KARYOMORPHOLOGY OF FIVE TAXA OF ZEPHYRANTHES HERB. FROM SOUTH INDIA

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INTRODUCTION

Zephyranthes Herb., a genus of about 65 species of bulbous flowering plants which are natives of warmer part of America and of West Indies belongs to the family Amaryllidaceae (Hutchinson, 1973, Judd et al., 1999). Many of them are valued as ornamentals with relatively small flowers and short life cycle (Lorenzi and Souza 1999). The leaves are narrowly strap shaped and few in number, flowers solitary and are usually borne on a pedicel of variable length. Cytology of over 50 species have been investigated (Darlington and Wylie 1955, Fedorov 1969 and Goldblatt 1981, 1984, 1985 and 1988), which are taxonomically complex and cytologically variable group. However, the quest for size and colour of flower has led to indiscriminate use of hybrids and wild species in breeding and the evolution of cultivars warrants their screening. This has prompted the present authors to take up this study. Present study deals with detailed karyotype analysis of five taxa.

MATERIALS AND METHODS

Plant specimens were collected from different localities of South Indian states of Kerala and Tamil Nadu. For cytological analysis, young root tips were pre-treated in 0.002 M aqueous solution of 8-hydroxyl-quinoline for 3 hrs at 40 C and then fixed in Carnoy's fluid for 24 hrs and stored in the refrigerator. The roots were first washed twice for 5 min in distilled water, hydrolyzed for 20 min in 5N HCl and then squashed. The chromosomes were stained in 2% aceto carmine. Photomicrographs were taken from temporary preparations. Karyomorphological analysis was made following Stebbins (1958) and Levan et al (1964). Chromosomes were arranged in three groups according to the relative position of the centromere (metacentric-m, centromeric index-Ci= 50 to 37.5, submetacentric-sm, Ci= 37.5 to 25 and subtelocentric-st, Ci=25 to 12.5 based on arm ratio and centromeric index (Stebbins, 1958). Cells were photographed with Olympus photomicroscope. After species identification, voucher material was deposited at Herbarium of University of Kerala, Botany Department.

RESULTS AND DISCUSSION

The karyotype details of the five taxa studied are given in (Table 1) and idiograms in (Fig. 2). Z.grandiflora revealed 2n=48, Z. longifolia 2n= 46, Z. candida 2n=25 and Z. rosea 2n=24, while the four accessions of Z. mesochola showed 2n=36, 19, 18 and 16 chromosomes each respectively (Fig. 1).

The presently studied five taxa of *Zephyranthes* from South India are all based on x=6. The karyotype of all the species are graded with five of them belonging to the 2B category and three of them to 3B category. Z. candida, a tetraploid based on x=6 was less specialized with mostly m-type and sm-type of chromosomes. Chromosome number 5 in Z. candida was heteromorphic with only one of the members of homologue possessing the satellite while in chromosome number 7 both members possessed satellite. Z. rosea is a tetraploid on x=6 and showed more asymmetry with 2- st, 16- sm, and 6-m type of chromosomes. The chromosome 3 possessed satellite. The hexaploid Z. longifolia was less specialized with mostly m-type and sm-type of chromosomes. Z. grandiflora representing the octaploid based on x=6, revealed a graded karyotype which belonged to the category 2B with m- and sm- type of chromosomes. Of the four taxa of Z. mesochola, variability in chromosome number were evident, there was little karyomorphological change with the different taxa having median, submedian and subterminal constrictions. It is apparent that apart from the superficial similarity in chromosome morphology of the different species in having long, medium as well as short chromosome with either median, submedian and subterminal constrictions, they differ from each other in their chromosome complement, total amount of chromatin matter, average chromosome length as well as TF%. On the whole, the karyomorphological situation in the genus Zephyranthes reveal that structural alterations of chromosomes to be less operative. However, chromosome numerical data holds the genus to constitute a complex with an array of chromosome numbers ranging from 10 to 200 with 12, 18, 22, 24, 26, 28, 40-50, 54, 58-61, 66-69, 72, 73, 93, 100, 120 and 200 of which most frequent number being 2n=48 (Inariyama 1937, Coe 1954, Yokouchi 1963, Flory and Smith, 1980a and Davina, 2001). Also 2n=24 and 36 (Kapoor and Tandon 1963), 2n=38 (Nagao and Takusagawa 1932), 2n=40 (Tandon and Meenakshi, 1965) have been reported. Some authors considered x=6 to be the basic number for the genus (Inariyama 1937, Sato 1938, Sharma and Ghosh 1954, Flagg 1960 and Raina and Khoshoo 1971). Darlington and Wylie (1955) suggested three basic numbers of x=6, 7 and 9 for this genus. However x=6 seems to be deep rooted basic constitution of the entire tribe Zephyrantheae. Sato (1938) has opined that x=6 condition may have originated from x=11 by fusion of translocation or by loss of one pair of chromosome from a 7-chromosomal ancestor.

Chromosome number variability in 32 individuals of Z. sylvatica populatopn from Northeast Brazil was investigated by Felix et al. (2008). They identified three cytotypes-2n= 12 (one metacentric, four submetacentric and one acrocentric pairs) in 24 individuals; 2n=12+1B in five and three individuals with 2n=18, a triploid cytotype. Here all diploid individuals showed chromosomes with polymorphism in pair one and two. But in triploids the polymorphism was observed in all chromosome triplets with two homomorphic

chromosomes and a higher or lower heteromorphic chromosome. All species showed karyotypes formed by a set of metacentric to submetacentric chromosomes and a smaller number of acrocentric chromosomes, with gradual reduction in size and chromosome morphology. Different species had distinct karyotypic formula and sometimes they coincided concerning ploidy level (Felix et al., 2011).

In this context it is pertinent to note that all the varying nuclei show their chromosome complement as multiples of six and this irregularity has been explained on the basis of endomitotic replication (Sharma and Ghosh 1956). Thus the polyploid species of Zephyranthes occur in a regular series in multiples of x=6 ranging from 2x to 16x showing a great role of polyploid evolution on this number in this group.

The chromosome in this family is large and it is apparent that considerable amount of segmental interchange coupled with aneuploidy and polyploidy, due probably to prolific vegetative multiplication have enabled these forms to multiply. Further evolution as it is evident must have been through aneuploidy in both directions at different ploidy levels giving rise to taxa with a variety of chromosome constitutions. Of the five taxa presently studied all of them are based on x=6, one of each with 2n=48 (Z. grandiflora), 2n=36 (Z. mesochola), 2n=24 (Z. rosea) and the rest are aneuploids with 16, 19, 25 and Z. longifolia with 2n=46 respectively. The overall chromosome situation appears to be suggestive that this genus is an evolutionarily dynamic one and that chromosomal numerical variation, both polyploidy and aneuploidy, have contributed significantly more than structural rearrangement in the rapid evolution of this genus.

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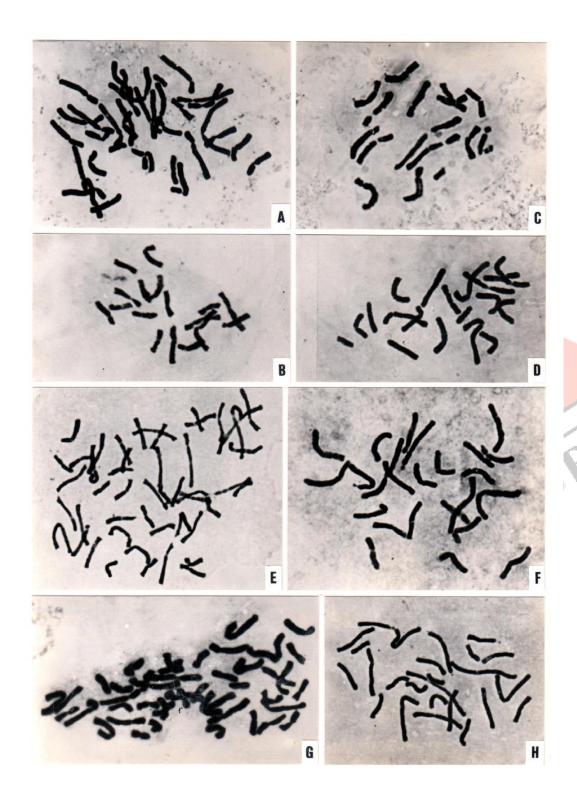


Fig. 1 (A-H) - Somatic chromosome of *Zephyranthes* (x900).

A. Z. mesochola, accession IV (2n=36), B. Z. mesochola, accession II (2n-2=16), C. Z. mesochola, accession III (2n=18), D. Z. mesochola, accession I (2n+1=19), E. Z.

grandiflora (2n=48), F. Z. rosea (2n=24), G. Z. longifolia (2n-2=46), H. Z. candida (2n+1=25).

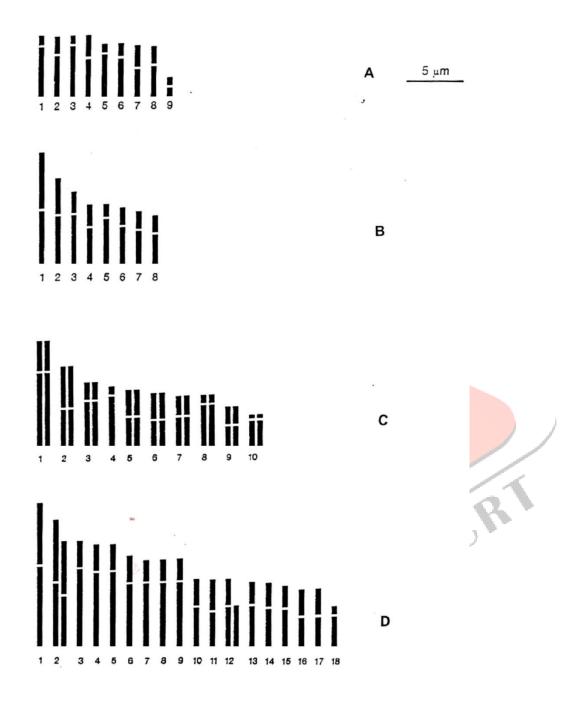


Fig. 2. (A-H) - Idiograms of the haploid complements of Zephyranthes.

A. Z. mesochola, accession III, B. Z. mesochola, accession II, C. Z. mesochola, accession I, D. Z. mesochola, accession IV..

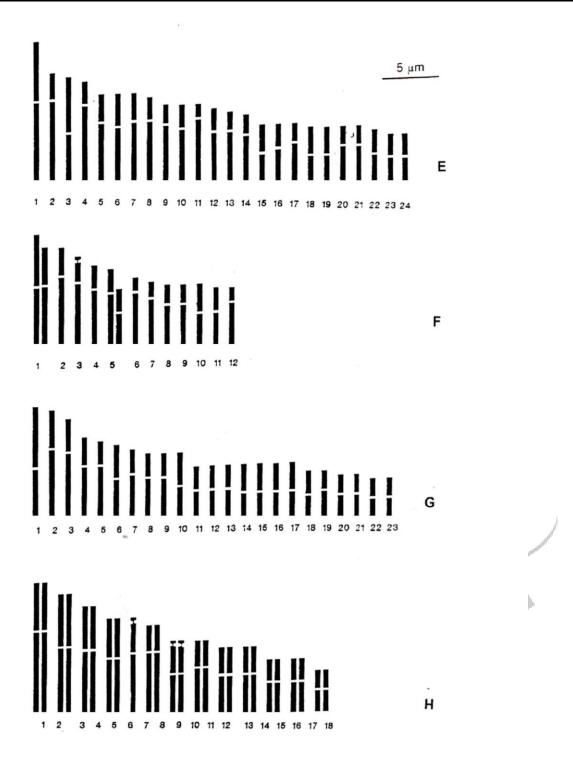


Fig. 2. (A-H) - Idiograms of the haploid complements of Zephyranthes.

E. Z. grandiflora, F. Z. rosea, G. Z. longifolia, H. Z. candida.

TABLE 1. KARYOMORPHOLOGICAL DATA ON EIGHT TAXA OF ZEPHYRANTHES FROM SOUTH INDIA

| Taxon | Karyotype formula | TCL (µm) | ACL (µm) | TF% | Karyotype category (STEBBINS 1958) |
|---------------------------|-------------------------------------------------------------------------------------------|-------------|-------------|-------|------------------------------------------|
| Z candida | 2n = 25: 2 Bsm + 2 Bm + 2 Cm + 5 Dsm + 2 Em + 2 Fsm + 4 Fm + 6 Gm | 237.73 | 9.50 | 37.21 | 2B |
| Z rosea | 2n = 24; 1 Cm + 3 Dsm + 5 Esm + 1 Fm + 2 Fst + 6 Fsm + 4 Fm + 2 Gsm | 191.01 | 7.95 | 31.38 | 38 |
| Z longifolia | 2n = 46; 2 Cm + 4 Csm + 2 Em + 6 Esm + 6 Fm + 4 Fsm + 18 Gm + 4 Hm | 323.02 | 7.02 | 40.80 | 2B |
| Z grandiflora | 2n = 48: 2 Am + 2 Cst + 2Cm + 6 Dsm + 2 Dst + 6 Esm + 4 Est + 6 Fsm + 6 Fm + 8 Gm + 4 Gsm | 410.42 | 8.55 | 32.76 | 3B |
| Z mesochloa accession l | 2n = 19; 2 Csm + 2 Em + 2 Esm + 1 Fst + 4 Fm + 4 Gm + 2 Gst + 2 Hst | 136.82 | 7.20 | 33.07 | 2B |
| Z mesochloa accession II' | 2n = 16; 2 Cm + 2 Dm + 2 Esm + 6 Fsm + 2 Fst + 2 Gsm | 127.42 | 7.96 | 36.25 | 2B |
| Z mesochloa accession III | 2n = 18; 2 Fm + 6 Fsm + 6 Fst + 2 Gm + 2 Hm | 109.28 | 6.07 | 28.98 | 38 |
| Z mesochloa accession IV | 2n = 36: 2 Am + 1 Bm + 1 Cm + 6 Cst + 2 Dsm + 6 Dst + 2 Em + 7 Fm + 6 Fsm + 3 Gst | 352.58 | 9.79 | 33.01 | 2B |

TCL – Total chromosome length of diploid complement; ACL – Average chromosome length; TF% - (Total short arm length \times 100) / (Total long arm length \times Total short arm length).

