APPENDIX

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ESTIMATION OF DNA OF SOME TAXA INVESTIGATED

During the past several years, abundant data have been obtained regarding deoxyribonucleic acid (DNA) content per nucleus and per chromosome among plant species (Sato, 1963; Rothfels et al., 1966; Rees and Jones, 1967, 1972; Rothfels and Heimberger, 1968; Jones and Rees, 1968; Rees, 1972; Bennett, 1972; EL-Lakany, 1972; EL-Lakany and Dugle, 1972; Sparrow et al., 1972; SZ-Borsos, 1973; Nagl et al., 1973; Grant, 1969, 1976). Most of these works have been reviewed recently by Price (1976). A correlation of DNA content with ploidy level and total chromatin matter made suggest that the DNA content per nucleus of somatic cell is constant in each species (Sato, 1963). This supports the so-called DNA-constant hypothesis. Thus there is some relationship between chromosome number and DNA content per nucleus. However, the extensive variation in DNA content among angiospermic families and among genera and species of the families are known. Such observations have been made by Rees and Jones (1967), Bennett (1972), Sparrow and Neumann (1974) in the family Poaceae and by Rees and Hazarika (1967), Martin (1968), Chooi (1971), Bennett (1972) in the family Leguminosae.

Though there is no overall correlation between chromosome

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size or nuclear DNA content and phylogeny in angiosperms (Stebbins, 1938, 1950, 1966), the distribution of DNA content is very specific. Studies of closely related plant species indicate that both evolutionary increase and decrease in DNA content is apparently common. Rees and Hazarika (1967) based on determination of the nuclear DNA content of eighteen diploid (2n = 18) species of <u>Lathyrus</u> have proposed that, diminution of DNA content must have occurred during evolution of species. Similarly Nagl <u>et al.</u>,(1973) while working with family Asteraceae have shown that, the phylogenetic reduction in chromosome size is closely related with reduced DNA-content of a diploid nucleus.

In the present study an attempt is made to see whether a similar correlation (i.e. correlation of reduced DNA-content with phylogenetic reduction) exists in the taxa investigated. Totally 20 species belonging to the genera <u>Indigofera</u>, <u>Desmodium</u> and <u>Alysicarpus</u> of Fabaceae are selected for the present work.

MATERIALS AND METHODS :

Seedlings at open cotyledon stage (7 days old) were taken for the estimation. The different species for which estimation is carried out are listed in Table 1. ñ

Preparation of extract :

A 20% homogenate of the radicle free seedlings was prepared by grinding in pestle and morter for 20 min. in methanol at room temperature. This homogenate was used for the estimation of DNA following Schneider's method (1957).

Estimation of DNA :

The determination of DNA is based on its preferential solubility in hot trichloro-acetic acid (TCA), which is quantitated by means of colorimetric reactions involving the pentose component of DNA.

5 ml of homogenate was centrifuged at 1500 x g for 10 min. and the residue obtained was made pigment free by giving 5-6 washes following the method of Smille and Krotkov (1960). The residue was then suspended in 2 ml of 5% TCA and allowed to stand for 15 min. at room temperature. This was then centrifuged for 15 min. at 1500 x g. The residue obtained was suspended in 5 ml of 95% ethanol, kept in boiling water for 30 seconds, cooled and centrifuged at 1500 x g for 15 min. The residue obtained was suspended in 2 ml of 5% TCA, kept in hot water both at 90°C for 15 min. by occasional stirring. This was cooled and centrifuged at 1500 x g for 15 min., the supernatant was collected. The residue was washed with Fig. 1 A - Standard DNA graph.

Fig. 1 B - Comparision of total chromatin length with total DNA-content in :

1. Indigofera astragalina

3. <u>I. trifoliata</u>

5. <u>I. tinctoria</u>

7. <u>I.linifolia</u> var.<u>campbelli</u> 8. <u>I. linifolia</u>

9. <u>I. oblongifolia</u>

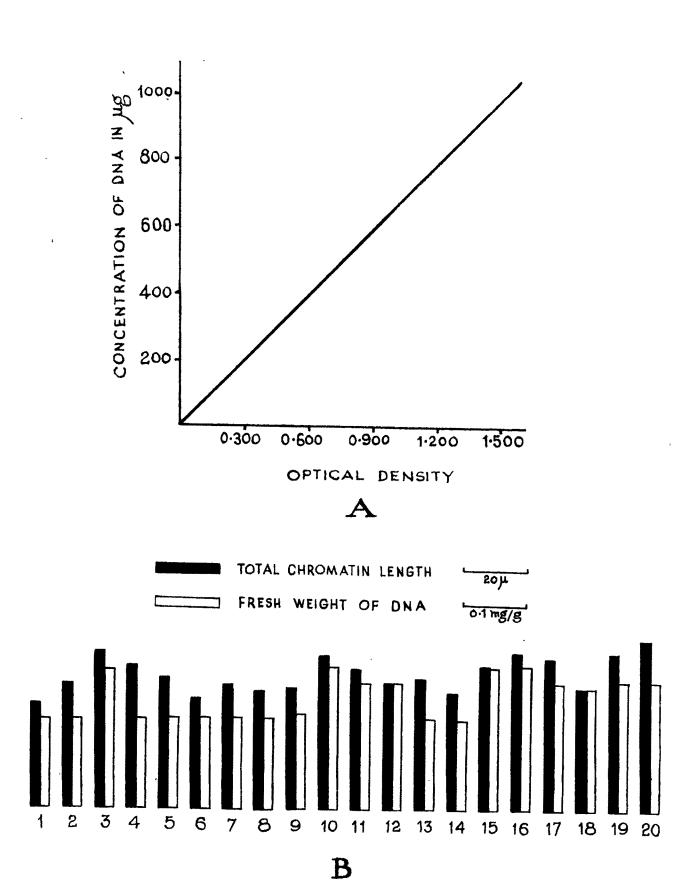
11. Alysicarpus bupleurifolius

- 13. <u>A. longifolius</u>
- 15. <u>A. wallichii</u>

17. <u>A. monilifer</u>

19. Desmodium dichotomum

- 2. I. hirsuta
- 4. <u>I</u>. <u>linnaei</u>
- 6. <u>I. cordifolia</u>
- 10. I. glandulosa
- 12. A. procumbens
- 14. <u>A. heyneanus</u>
- 16. <u>A. styracifolius</u>
- 18. A. vaginalis
- 20. D. gangeticum



2 ml 5% TCA and the supernatant obtained was added to the previous collection. This was used for DNA estimation using Diphenylamine reagent and the blue colour developed was read at 660 nm. using Carl Zeiss Spectrophotometer. Purified Calf Thymus DNA (BDH) was used as standard (Fig. 1A).

RESULTS AND DISCUSSION :

The results are summarised in the Table 1. As can be seen from the table that DNA concentration ranges from 0.044% to 0.061% by dry weight. Comparision of total DNA content with total chromatin length of the species investigated show good correlation (Fig. 1B). The increase in DNA content with the increase of total chromatin length is observed in majority of the species (Fig. 1B). Rees and Hazarika (1967) have shown , diminution of DNA content with the advancement of the species in the genus <u>Lathyrus</u>. Nagl <u>et al</u>., (1973) while, working with Asteraceae have also shown that, the phylogenetic reduction in chromosome size is closely related with reduced DNA content of diploid nucleii. The species of the genera <u>Indigofera</u>, <u>Desmodium</u> and <u>Alysicarpus</u> investigated presently do not show such correlation.

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V	· · · · · · · · · · · · · · · · · · ·		~	Ň		Residue	,
	, ,	۰. ۲	· · ·	ζ. ·	Residue 5% TCA 90 ⁰ C (Twice)	-	
			, , , ,	Residue 95% Ethanol boiling water bath	Re 5% TCA	Supernatant (DNA)	
		due with methanol	Residue 5% TCA		Supernatant	Ň	•
• •	Tissue (homogenate in methanol)	Residue Washed 5 times with r	Supernatant	Supernatant		2	
Λ	Tissue in m	Supernatant	ທີ `		,	, . ,	•

DNA Total Somatic Chromatin Sr. Coll. mg/g g/100 g Taxa Number length No. No. fresh wt. dry wt. (2 n) (in µ) 1. Indigofera astragalina 6. 32.56 0.182 16 0.044 0.045 2. I. hirsuta 102 16 38.76 0.185 30 48.58 0.233 0.057 3. I. trifoliata 16 4. I. linnaei 23 16 . 44.14 0.182 0.044 0.044 0.182 5. I. tinctoria 4 16 40.60 6. <u>I. cordifolia</u> 25 16 34.42 0.185 0.045 7. I.linifolia var. campbelli 18 38.66 0,182 0.044 16 0.045 16 36.78 0.185 8. <u>I. linifolia</u> 19 0.048 9. I. oblongifolia 0.195 16 37.10 11 0.059 47.96 0.242 10. I. glandulosa 16 16 . 0.196 0.049 11. Alysicarpus bupleurifolius 101 16 43.48 0.194 0,047 16 ,39.18 12. A. procumbens 83 0.045 40.24 0.182 16 13. A. longifolius 82 0.185 01045 36.20 34 16 14. A. heyneanus 0.242 0.061 44.52 89 16 15. A. wallichii 0.059 16. A. styracifolius 40 16 48.46 0.249 0.192 0.046 36.96 39 16 17. A. monilifer 0.044 36.96 0.182 18. A. vaginalis 66 16 0.050 48.30 0,209 19. Desmodium dichotomum 22 50 0.205 0.050 20. D. gangeticum 52.60 49 22

Table 1. Comparision of the total chromatin length with total DNA content of the taxa investigated.