Nuclear DNA Variation in Relation to Cytological Features of Some Species in the Genus *Plantago* L.

A. Badr¹, R. Labani² and T. T. Elkington³

Accepted February 20, 1986

During the last 15 years or so abundant data have been accumulated indicating that a wide variation exists among plants in the nuclear DNA content (Rees 1972, Rees and Jones 1972, Price 1976, Bennet et al. 1982). Several attempts have been made to utilise this variations in the study of species relationship in various taxonomic groups (Sparrow and Nauman 1974, Price and Bachman 1975, Greilhuber 1977). These studies have shown that variation in nuclear DNA content often provides useful evidence which helps to assess the relationships among related species particularly when used in combination with other features particularly cytological characters.

The genus *Plantago* L. (Plantaginaceae) was monographed by Pilger (1937) and was revised by Rahn (1978) who proposed some nomenclatorial changes and suggested a key for the subdivisions and some sections of the genus. The cytological studies carried out on *Plantago* indicate that about 70% of the species in this genus have a basic chromosome number of \( x=6 \) and are distributed in the three subgenera of the genus. A basic number of \( x=5 \) was recorded in about 25% of the species in *Plantago* which are placed in subgenus *Psyllium* (Juss.). Harms and Reiche or in subgenus *Coronopus* (Lam and DC) Rahn. A basic number of \( x=4 \) is so far only recorded in two species comprising less than 2% of the genus and they are placed in subgenus *Psyllium* (McCullagh 1934, Rahn 1957, Fernandes and Franca 1973, Zemskova 1977).

In Egypt the genus *Plantago* comprises 21 species. The majority of which are found in the coastal Mediterranean region. In the present study the nuclear DNA in 9 species of the genus are measured and related to the cytological features and taxonomic relationships of the studied taxa.

Material and methods

Material of the studied species was collected from their natural habitats from various parts of Egypt. A list of these species and localities from which they were collected is given in Table 1. The distribution of localities in the different parts of Egypt is shown on the map (Fig. 1). Identification of species was made by the senior author (A. Badr) and verified by Prof. M. N. El-Hadidi, Professor of Plant Taxonomy and Head of Botany Department, Cairo University. Vouchers are kept at the herbarium of Botany Department, Tanta University.

For the determination of DNA per nucleus seeds were germinated on moist filter paper simultaneously with seeds of *Allium cepa* cv. Ailsa Craig which was used as a standard in the determination of the \( 4C \) DNA amounts. Root tips of samples and that of *A. cepa* were fixed in 3:1 alcohol: glacial acetic acid for 24 hours and were then washed in distilled water, hydrolized in 1N HCl at 60°C for 10 minutes and stained in leucobasic fuchsin for 2 hours. Roots were then washed in running water for 8 minutes and in \( \text{SO}_2 \) water for 30 minutes and stored in distilled water. Squash preparations were made in a drop of distilled water and coverslips

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¹ Botany Department, Faculty of Science, Tanta University, Egypt.
² Botany Department, Al-Fateh University, Tripoli, Libya.
³ Botany Department, Sheffield University, England.
were separated by quick freezing in carbon dioxide ice. Preparations were dehydrated in alcohol and mounted in euparal.

DNA measurements were carried out using a Vickers M86 scanning microdensitometer with a range of different mask sizes depending on the area of the nucleus. Measurements were made on 10 prophase nuclei per root tip and 3 root tips for each sample were used. Comparisons of the mean of these measurements with that of A. cepa were made and the 4C DNA amount of each species of Plantago was calculated using the following formula: 

\[ 4C \text{ DNA} = \frac{A}{B} \times C \]

where A = the 4C DNA amount in A. cepa derived from 2C determinations (Bennet et al. 1982), B = the mean absorption figure measured in the A. cepa standard root tips and C = the mean absorption figure measured in a species of Plantago.

Chromosome numbers and mean chromosome length of the studied species were determined in root tips pretreated with 0.05% colchicine solution and fixed in 3:1 alcohol: acetic acid. Preparations were made using the Feulgen squash method and were made permanent by mounting in canada balsam. The mean 4C DNA value recorded was divided by the basic number of chromosomes to obtain the DNA/genome so that data of the different samples can be compared.

Results and discussion

The amount of DNA and cytological features of the 14 samples belonging to 9 species of Plantago are given in Table 1. The two samples of P. albicans L. contain the highest 4C DNA amount/nucleus among the studied species; being 7.77 pg in sample 1 and 7.37 pg in sample 2 respectively. Both samples are hexaploid with 2n = 30, but the karyotype of sample 2 also includes 3B chromosomes. In the tetraploid P. crassifolia Forsk. (2n = 20) the recorded 4C DNA was 5.87 pg. In P. crypsoides Boiss. and the two samples of P. cylindrica Forsk. which are diploid (2n = 10) but resemble the two previous species in having a basic chromosome number of x = 5, similar amounts of 4C DNA/nucleus were measured (Table 1).

The measured 4C DNA content/nucleus in two samples of P. lagopus L. and in P. lanceolata L. both diploid with 2n = 12 and x = 6 is similar, but considerably higher than those recorded in the diploid samples with x = 5 i.e. P. crypsoides and P. cylindrica (Table 1). The 4C DNA content/nucleus was measured in two other diploid species with x = 6. In P. major L. the measured 4C DNA amount/nucleus was 3.90 pg which is smaller than those recorded in P. lagopus and P. lanceolata. In P. notata Lag., on the other hand, the highest amount of 4C DNA amount/nucleus among the diploid species investigated (5.73 pg) was recorded (see Table 1).

The 4C nuclear DNA was measured in 3 samples of P. ovata Forsk. The two diploid samples (1 and 2) both with 2n = 8 have 2.14 pg and 1.98 pg DNA in the 4C nucleus respectively. In the tetraploid sample of this species (2n = 16) a DNA amount of 3.2 pg in the 4C nucleus was recorded.

The amount of DNA per genome in the two hexaploid samples of P. albicans and the tetraploid species P. carssifolia was found to be smaller than the corresponding amounts in the diploid species having the same basic number (x = 5) i.e. P. crypsoides and P. cylindrica (Table 1). Similarly the DNA amount-genome in the tetraploid sample of P. ovata (x = 4) was con-
The DNA amount/genome in the species with \( x=6 \) was remarkably higher than those found in species with \( x=5 \) or \( x=4 \). Thus, the amount of DNA/genome is generally correlated with the basic chromosome number of the *Plantago* species studied. *P. ovata* with \( x=4 \) contains the least amount of DNA/genome. Higher amounts of DNA/genome were found in the species with \( x=5 \) i.e. *P. albicans*, *P. crassifolia*, *P. crypsoides* and *P. cylindrica*. However, the amounts of DNA/genome recorded in the species having \( x=6 \) i.e. *P. lagopus*, *P. lanceolata*, *P. major* and *notata* were higher than those found in the species with \( x=5 \) (Table 1).

It is also evident from the present data (Table 1) that the amounts of DNA/nucleus in the polyploid species i.e. *P. albicans* (2n=30, \( x=5 \)), *P. crassifolia* (2n=20, \( x=5 \)) and sample of 3 of *P. ovata* (2n=16, \( x=4 \)) have much higher DNA content compared with diploid species having the same basic number. These findings support the conclusion of Price (1976) that polyploidy may be accompanied by the duplication of nuclear DNA. However, the increase in nuclear DNA in the polyploid species is lower than expected on the basis of the ratio between the diploid and hexaploid level (1:3) as in *P. albicans* or the tetraploid level (1:2) as in *P. crassifolia* or *P. ovata*. The ratio between the nuclear DNA in *P. crypsoides* (2n=10, \( x=5 \)) and the average amount of the two samples of *P. cylindrica* (2n=10, \( x=5 \)) and that in *P. albicans* are 1:2.06 and 1:2.17 respectively. The ratio between the nuclear DNA in these two species and that found in *P. crassifolia* are 1:1.6 for *P. crypsoides* and 1:1.68 for *P. cylindrica*. The ratio between the average nuclear DNA of the two diploid samples of *P. ovata* and the tetraploid sample of the same species is 1:1.55. Similar results were found among diploid and hexaploid plants of *Betula* (Taper and Grant, 1973). These authors reported a ratio of 1:2.19 between diploid (2n=28) and hexaploid (2n=84) plants of *Betula*.

**P. albicans** and **P. cylindrica** are taxonomically grouped together in subgenus *Psyllium* (Juss.) Harms & Reiche and also in the same section *Albicans* Barns and the same series *Albicantes* (Rahn 1978). *P. ovata* is also placed in subgenus *Psyllium* section *Albicans* but in series *Ovatae*. Thus the similarities in the DNA amount/genome between these species support their taxonomic affinities based on morphological criteria. *P. crassifolia*, on the other hand, is placed in subgenus *Coronopus* (Rahn 1978), whereas *P. crypsoides* is morphologically different.

### Table 1. Localities, cytological features and DNA contents per nucleus (4C DNA), per genome in the studied species

<table>
<thead>
<tr>
<th>Species</th>
<th>Locality</th>
<th>Chrom. number (2n)</th>
<th>Basic number (x)</th>
<th>Ploidy level</th>
<th>4C DNA</th>
<th>DNA/ genome</th>
<th>Mean chrom. length in ( \mu )m</th>
</tr>
</thead>
<tbody>
<tr>
<td>1. <em>P. albicans</em> 1</td>
<td>Burg El-Arab</td>
<td>30</td>
<td>5</td>
<td>6x</td>
<td>7.77</td>
<td>1.29</td>
<td>3.44</td>
</tr>
<tr>
<td>2. <em>P. albicans</em> 2</td>
<td>Sinai, El-Arish</td>
<td>30+3B</td>
<td>5</td>
<td>6x</td>
<td>7.37</td>
<td>1.23</td>
<td>3.62</td>
</tr>
<tr>
<td>3. <em>P. crassifolia</em></td>
<td>25 km West Alex.</td>
<td>20</td>
<td>5</td>
<td>4x</td>
<td>5.87</td>
<td>1.47</td>
<td>2.90</td>
</tr>
<tr>
<td>4. <em>P. crypsoides</em></td>
<td>Burg El-Arab</td>
<td>10</td>
<td>5</td>
<td>2x</td>
<td>3.68</td>
<td>1.88</td>
<td>2.55</td>
</tr>
<tr>
<td>5. <em>P. cylindrica</em> 1</td>
<td>140 km South Alex,</td>
<td>10</td>
<td>5</td>
<td>2x</td>
<td>3.30</td>
<td>1.65</td>
<td>3.03</td>
</tr>
<tr>
<td>6. <em>P. cylindrica</em> 2</td>
<td>Sinai, Sidr</td>
<td>10</td>
<td>5</td>
<td>2x</td>
<td>3.84</td>
<td>1.97</td>
<td>3.13</td>
</tr>
<tr>
<td>7. <em>P. lagopus</em> 1</td>
<td>Burg El-Arab</td>
<td>12</td>
<td>6</td>
<td>2x</td>
<td>4.61</td>
<td>2.31</td>
<td>2.40</td>
</tr>
<tr>
<td>8. <em>P. lagopus</em> 2</td>
<td>Madinet Qasr, Cairo</td>
<td>12</td>
<td>6</td>
<td>2x</td>
<td>5.00</td>
<td>2.50</td>
<td>2.62</td>
</tr>
<tr>
<td>9. <em>P. lanceolata</em></td>
<td>25 km South Alex.</td>
<td>12</td>
<td>6</td>
<td>2x</td>
<td>5.16</td>
<td>2.58</td>
<td>2.96</td>
</tr>
<tr>
<td>10. <em>P. major</em></td>
<td>Kafr El-Sheikh</td>
<td>12</td>
<td>6</td>
<td>2x</td>
<td>3.82</td>
<td>1.91</td>
<td>2.04</td>
</tr>
<tr>
<td>11. <em>P. notata</em></td>
<td>Burg El-Arab</td>
<td>12</td>
<td>6</td>
<td>2x</td>
<td>5.73</td>
<td>2.87</td>
<td>4.01</td>
</tr>
<tr>
<td>12. <em>P. ovata</em> 1</td>
<td>Sallum</td>
<td>8</td>
<td>4</td>
<td>2x</td>
<td>2.14</td>
<td>1.07</td>
<td>3.04</td>
</tr>
<tr>
<td>13. <em>P. ovata</em> 2</td>
<td>50 km East Cairo</td>
<td>8</td>
<td>4</td>
<td>2x</td>
<td>1.98</td>
<td>0.99</td>
<td>3.13</td>
</tr>
<tr>
<td>14. <em>P. ovata</em> 3</td>
<td>Madinet Nasr, Cairo</td>
<td>16</td>
<td>4</td>
<td>4x</td>
<td>3.20</td>
<td>0.80</td>
<td>3.22</td>
</tr>
</tbody>
</table>
from *P. albicans* and *P. cylindrica*. The sectional delimitation of *P. crypsoides*, however, needs further investigation.

*P. lagopus* and *P. lanceolata* share several morphological and cytological criteria and also have similar amounts of nuclear DNA. *P. lanceolata* is placed in subgenus *Psyllium* and is considered to be the type species of section *Lanceifolia* Barns. In view of the similarities between these two species *P. lagopus* may be placed in section *Lanceifolia* of subgenus *Psyllium*. *P. notata* contains a nuclear DNA amount which is to some extent similar to that recorded in *P. lanceolata*. *P. notata*, however, is cytologically characterized by having longer chromosomes when compared with other species with *x*=6 and is morphologically distinct from them by its penatifid leaves and wooly spikes. The sectional delimitation of this species is yet to be made.

*P. major* also with 2n=12 and *x*=6, on the other hand, contains smaller amount of nuclear DNA if compared with other species having *x*=6. Cytologically this species is characterized by short chromosomes and symmetric karyotype. Morphologically *P. major* is also distinct by its 3–9 veined, broad, ovate, glabrous leaves, narrow, green, long spikes and its affinity to grow in moist places. Taxonomically this species is separated from the other studied species and is placed in subgenus *Plantago* by both Pilger (1937) and Rahn (1978). Based on the present results and the morphological and cytological criteria the separation of *P. major* seems justified.

The basic chromosome number of *x*=6 is the most common among the species of *Plantago* and may be considered the ancestral number in the genus (Setbbins and Day 1967). The basic number of *x*=4 may represent the end point of a stepwise reduction in the basic number. A similar pattern of change in the nuclear DNA may be deduced from the present results. Species with *x*=6 have considerably higher amounts of nuclear DNA in comparison to those with *x*=5 or *x*=4, whereas species with *x*=4 have the least amount of nuclear DNA. It is therefore suggested that the reduction in chromosome number has been associated with a reduction in the amount of nuclear DNA.

**Summary**

The nuclear DNA content in 14 taxa belonging to 9 species of *Plantago* from Egypt has been measured using cytophotometric methods. The recorded DNA amounts were related to the karyotype features of the studied species. In general the amount of nuclear DNA-genome is correlated with the basic chromosome number. Species having *x*=6 i.e. *P. lagopus*, *P. lanceolata*, *P. major* and *P. notata* contain higher amounts of nuclear DNA-genome when compared with the species having *x*=5 i.e. *P. albicans*, *P. crassifolia*, *P. crypsoides* and *P. cylindrica*. Three samples of *P. ovata* (*x*=4) were found to contain the least amounts of DNA-genome. It was also found that polyploid taxa contain higher amounts of 4C DNA/nucleus than diploid species. The impact of these results on the taxonomic relationships of the studied species is discussed.

**References**


