Karyological and morphological variations within the genus Dysphania (Chenopodiaceae) in Bulgaria

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Abstract – The karyological and morphological variability of species from the genus Dysphania were studied. The results demonstrated that genus Dysphania is represented in Bulgaria by five species: Dysphania ambrosioides, D. multifida, D. botrys, D. schraderiana and D. pumilio. The first two species are tetraploids with chromosome number $2n = 32$ for D. ambrosioides and $2n = 36$ for D. multifida. The remaining three species are diploids with $2n = 18$. The results from statistical analysis demonstrated that the main source of phenotype variation in the species is the interpopulation variation. The specific characters which allowed their recognition are the morphological characteristics of the perianth lobes, the upper leaves and the seeds. The distinction between D. multifida, D. ambrosioides and D. schraderiana is based on differences in the quantitative traits, while in D. botrys and D. pumilio qualitative traits are also important. The basic evolutionary mechanisms are polyploidy and diploidy. A tendency towards reduction in the size of generative organs and the number of perianth lobes was found.

Key words: Dysphania, distribution, ecology, karyology, morphology, Bulgaria

Introduction

Traditionally, the genus Dysphania comprises 6–10 species, endemic for Australia (MOSYAKIN 1993, CLEMENTS and MOSYAKIN 2003, ZHU et al. 2003). Over the past decades, the systemic definition and composition of the genus has frequently been a subject of discussion (MOSYAKIN 1993; MOSYAKIN and CLEMENTS 1996, 2002, 2008). Currently the genus Dysphania comprises about 40 species that earlier constituted the subgenus Ambrosia of the genus Chenopodium or were referred to the genera: Neobotritium Moldenke, Rouhieva Moquin-Tandon, Teloxys Moquin-Tandon. The representatives of the genus are mainly ruderal and weed plants, more common in the tropics, subtropics and warm-temperature...
Most of them entered Europe through the import of wool and agricultural products from Australia, South America, Eastern Asia, and Africa. A number of researchers (Aellen 1960, Uotila and Suominen 1976, Dostalek 1985, Kühn 1993, Uotila 2001) pointed out their great morphological variability, especially prominent in the leaf lamina. The similar, rarely exceeding 1 mm, dimensions of the generative organs very often make their distinction difficult.

For the Bulgarian flora so far five species have been reported – Chenopodium ambrosioides L. (= Dysphania ambrosioides (L.) Mosyakin et Clements), C. botrys L. (= Dysphania botrys (L.) Mosyakin et Clements), C. multifidum L. (= Dysphania multifida (L.) Mosyakin et Clements), C. pumilio R. Br. (= Dysphania pumilio (R. Br.) Mosyakin et Clements) and C. schraderianum Schult. (= Dysphania schraderiana (Schult.) Mosyakin et Clements) (Markova 1966, Assyov and Petrov 2006, Grozova 2007).

Dysphania ambrosioides forms populations on open sandy terrain, in close proximity to water basins from sea level to 250 m above sea level. It is usually a dominant species in anthropophyte or ruderal communities.

D. multifida forms populations on open lowland areas, more rarely on those facing west or south at altitudes from sea level to 700 m above sea level. It dominates or is an accompanying species in anthropophyte or ruderal communities dominated by cereal species.

D. botrys is widely spread on open, sunlit areas and as a weed in cultivated crops at altitudes from sea level to 900 m above sea level. Most often, it dominates or is an accompanying species in ruderal communities.

D. schraderiana was established for the Bulgarian flora by Pertti Uotila in 1993 while revising the materials from the genus Chenopodium in the herbarium of the Biological Faculty at Sofia University (SO). The species has very limited distribution. So far, only two populations have been known located in Vitosha and in Sredna Gora Mountain. Its ecological requirements are quite similar to those of D. botrys and specimens from both species were recorded in the population under study.


The objective of this study was to investigate the patterns and levels of morphological variation, chromosome numbers and karyotype morphology as well as ecological conditions of the natural local Bulgarian populations of genus Dysphania, and to attempt to trace the relationships between the species and some evolutionary trends in the genus.

Materials and methods

Morphological and karyological analyses were carried out on 27 natural Bulgarian populations of the genus Dysphania, referred to five species: Dysphania ambrosioides, D. botrys, D. multifida, D. pumilio, and D. schraderiana. Thirty plants from each population were used in the overall research.

Chromosome numbers and karyotypes have been reported on lasting preparations of metaphase root apex plates of seeds germinated in laboratory conditions collected in the nat-
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ural habitats of the species. The root tips were treated and squashed according to the accepted methods (GROZEV 2007). The chromosomal type was determined after the centromere index \( I = s/s+1 \), according to the classification proposed by GRIF and AGAPOV (1986). Three metaphase plates have been measured from each population. The voucher specimens are kept in the herbarium of the Bulgarian Academy of Sciences (SOM).


The mean value and coefficient of variation were calculated for each character of every population. They were used in the comparative analyses on different levels.

The relative contributions of intra- and interpopulation variation to the overall variation of each characteristic of the studied species were evaluated by one-way ANOVA.

Unweighted pair-group average (UPGA) hierarchical cluster analysis (HCA) was applied to the matrix with the Euclidean distances between the populations of the genus Dysphania in order to study the morphological pair-wise similarities and the underlying hierarchical classification structure.

The discriminant function analysis was conducted on morphological data to determine the most parsimonious way to distinguish between species. The stepwise discriminant analysis (SDA) procedure was used as it allows recognition of the most effectively discriminating variables among a large set of morphological characters.

Statistica 6.0 (StatSoft Inc. 2001) was used for the statistical analysis of morphological data.

Results

Karyology

As a result of karyological study, 3 chromosome numbers: \( 2n = 18 \), 32, 36 have been found in the genus Dysphania (Tab. 1). Two types of chromosomes: metacentric and sub-metacentric have been established in the karyotypes.

The number \( 2n = 18 \) has been counted in three species – Dysphania botrys, D. pumilio and D. schraderiana. The last species was studied karyologically in Bulgaria for the first time and for the studied population karyotype \( 2n = 14 \) m + 4 sm = 18 chromosomes has been established (Plate 1A). The diploid chromosome number found for D. schraderiana confirms the data known from literature sources (KJELLMARK 1934, SCHWARZOVA 1978a). For all three populations of D. pumilio the karyotype of \( 2n = 8 \) m + 10 sm = 18 chromosomes has been recorded (Plate 1B). The chromosome number \( 2n = 18 \) found for the Bulgarian popu-
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<td>Danubian plain, island Milka near Belene town, at 27 m, 43° 40’ N, 25° 10’ E, in sandy hair</td>
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<td>Thracian Lowland, Plovdiv town, near the river Maritza, at 164 m, 42° 09’ N, 24° 45’ E, in ruderal places</td>
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<td><em>Dysphania multifida</em> (L.) Mosyakin et Clemants</td>
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<td>No39 36</td>
<td>Danubian plain, Island Goljama Barzina near Belene town, at 20 m, 43° 40’ N, 25° 10’ E, in outskirts of the island and on sandy hair</td>
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<td>No40 36</td>
<td>Danubian plain, Vidin town, at 345 m, 43° 59’ N, 22° 52’ E, in ruderal places near the bus station with <em>Trifolium album</em> and other legumes.</td>
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<td>No105 36</td>
<td>Central Balkan Range, Kalofe town, at 666 m, 42° 37’ N, 24° 59’ E, grasslands with <em>Lolium perenne</em> L. of 50 m from post</td>
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<td>No106 36**</td>
<td>Western Sredna Gora Mt., Ikhtiman town, at 658 m, 42° 09’ N, 24° 45’ E, in ruderal places in the western parts of the city together with cereal species</td>
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<td>No107 36**</td>
<td>Eastern Rhodope Mts, Ivaylovgrad town, at 104 m, 41° 32’ N, 26° 08’ E, in ruderal places together with <em>Bromus tectorum</em> L. and <em>Setaria viridis</em></td>
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<td>Thracian Lowland, Parvomay town, at 134 m, 42° 06’ N, 25° 13’ E, in grasslands in kvartal Debar.</td>
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<tr>
<td>No109 36**</td>
<td>Thracian Lowland, Asenovgrad town, at 104 m, 42° 01’ N, 24° 52’ E, in ruderal places together with <em>Setaria viridis</em></td>
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| Sect. *Botryoides* (C. A. Mey) Mosyakin et Clements |
| *Dysphania botrys* (L.) Mosyakin et Clemants |
| No29 18 | Northern Black Sea Coast, around Varnensko ezero lake, at 79 m, 43° 11’ N, 27° 50’ E, periodically flooded coastal alluvial sands deposits |
| No97 18 | Southern Black Sea Coast, Pomorie town, at 0 m, 42° 35’ N, 27° 37’ E, in ruderal places near Monastery of Saint George |
| No100 18** | North-Eastern Bulgaria, Razgrad town, at 200 m, 43° 46’ N, 26° 31’ E, between the sidewalk and fences together with *Polygonum aviculare* |
| No35 18** | Danubian plain, Belene town, at 35 m, 43° 39’ N, 25° 07’ E, in ruderal places near the fishing pier |
| No102 18 | Eastern Balkan Range, Natural park »Sinite kamani«, at 290 m, 42° 42’ N, 26° 21’ E, in ruderal places near road from hotel Ablanovo to the lake Asenovets |
| No99 18** | Central Balkan Range, Trojan town, at 400 m, 43° 53’ N, 24° 43’ E, in ruderal places from 50 m to bus station |
KARYOLOGICAL AND MORPHOLOGICAL VARIATIONS IN *DYSPHANIA*

Tab. 1. – continued

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<td>Dysphania pumilio (R. Br.) Mosyakin et Clemants</td>
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<td>N°221 Thracian Lowland, Chirpan town, at 168 m, 42° 25′N, 25° 38′E, in ruderal places together with cereal plants</td>
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**Notes:**
- The karyotypes of *D. pumilio* correspond to the data of JAVOURKOVA-JAROMIROVA (1992) and MESIČEK (1992) from the Czech Republic, SCHWARZOVÁ (1978b) from Slovakia and RAHIMENIAD (2004) from Iran. With the third diploid species *D. botrys* differences have been recorded in the karyotype of the various populations. In population No. 29 from the Varna lake and population No. 30 from Plovdiv the karyotype consists of $2n = 4m + 14sm = 18$ chromosomes (Plate 1C). In population No. 99 from Troyan and population No. 45 from Tselina satellites can be seen on one pair of submetacentric chromosomes (Plate 1D). In the remaining eight populations (Tab. 1), a karyotype of 18 submetacentric chromosomes has been established. In four of these populations (No. 97 from Pomorie; No. 102 from »Sinite kamani« Natural Park; No. 53 from Gorno Novo selo; No. 100 from Razgrad) one of the chromosome pairs has satellites (Plate 2A), while in the other four (No. 35 from Belene; No.
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**Plate 1.** Microphotographs of the metaphase plate of *Dysphania* species: **A** – *D. schraderiana* (population No 103) – 2n = 18; **B** – *D. pumilio* (No 221) – 2n = 18; **C** – *D. botrys* (No 29) – 2n = 18; **D** – *D. botrys* (No 45) – 2n = 18. – Scale bars = 10 μm.

**Plate 2.** Microphotographs of the metaphase plate of *Dysphania* species: **A** – *D. botrys* (population No 53) – 2n = 18; **B** – *D. botrys* (No 98) – 2n = 18; **C** – *D. ambrosioides* (No 102) – 2n = 32; **D** – *D. ambrosioides* (No 201) – 2n = 32. – Scale bars = 10 μm.
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98 from Malo Buchino; No. 34 from Malka Vereya; No. 101 from Blagoevgrad) satellites have not been found (Plate 2B). The chromosome number $2n = 18$ registered for the Bulgarian populations of *D. botrys* confirms the familiar reference data (MULLIGAN 1961; MEHRA and MALIK 1963; KEENER 1970; BASSET and CROMPTON 1971, 1982; UOTILA 1973; QUEIROS 1975, SCHWARZOVÁ 1978b, 1980, 1993; DVOŘÁK et al. 1980; KHATOON and ALI 1993; LOMONOSOVA et al. 2003).

The tetraploid chromosome number $2n = 32$ has been found for *Dysphania ambrosioides*. The same result has been found in various regions in the geographical area of the species (LORZ 1937; WOROSCHILOV 1942; KAWATANI and OHNO 1950; RAGHAVAN and ARORA 1958; MEHRA and MALIK 1963; SHARMA and DEY 1967; MURIN and FERAKOVA 1974; QUEIROS 1975; SCHWARZOVÁ 1978b, 1986; SILVESTRE 1984; DVOŘÁK 1989; KHATOON and ALI 1993; AL-TURKI et al. 1999, 2000). For the three studied populations, differences were recorded in the karyotype structure. In the two populations from the Thracean lowland (No. 32 from Plovdiv and No. 104 from an island on the Maritsa River) the karyotype consists of $2n = 20 m + 12 sm = 32$ chromosomes (Plate 2C). For the population from the Danubian plain (No. 201 from the island of Milka) a karyotype of $2n = 4 m + 28 sm = 32$ chromosomes has been established (Plate 2D).

The highest chromosome number $2n = 36$ is registered in the populations of *D. multifida* (Tab. 1). The same number was reported by MARKOVA (1968) for another Bulgarian population of the species from the village of Koshava in the Danubian plain. The data from the other parts of its habitat are for $2n = 32$ (KAWATANI and OHNO 1956; GIUSTI 1970; QUEIROS 1975; GRANADO et al. 1988). In four of the studied populations (No. 39 from the island of Golyama Barzina, No. 108 from Parvomay, No. 105 from Kalofier, and No. 109 from Asenovgrad) the karyotype of $2n = 20 m + 16 sm = 36$ chromosomes has been registered (Plate 3A). In the other three populations (No. 40 from Vidin, No. 106 from Ihtiman, No. 107 from Ivaievgrad) the karyotype of $2n = 16 m + 20 sm = 36$ chromosomes has been established (Plate 3B).

![Plate 3](image)

Plate 3. Microphotographs of the metaphase plate of *Dysphania* species: **A** – *D. multifida* (population No. 105) – $2n = 36$; **B** – *D. multifida* (No. 40) – $2n = 36$. – Scale bars = 10 $\mu$m.

Morphology

The intrapopulation variation was estimated on the basis of the coefficient of variation (Tabs. 2–4). The data show that in all populations vegetative have higher level of variability than generative traits.
Tab. 2. Mean (first line) and coefficient of variation in % (second line) of *Dysphania schraderiana* (103), *D. pumilio* (214, 217, 221), *D. ambrosioides* (201, 32, 104) populations for each of the 24 observed characters

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KARYOLOGICAL AND MORPHOLOGICAL VARIATIONS IN *DYSPHANIA*

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GROZEVa N. H., CVETANOVA Y. G.

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</tbody>
</table>

Tab. 4. Mean (first line) and coefficient of variation in % (second line) of *Dysphania multifida* populations for each of the 24 observed characters

<table>
<thead>
<tr>
<th>Populations</th>
<th>103</th>
<th>214</th>
<th>217</th>
<th>221</th>
<th>201</th>
<th>32</th>
<th>104</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>51.2</td>
<td>90.9</td>
<td>73.4</td>
<td>88.6</td>
<td>83.6</td>
<td>54.2</td>
<td>65.5</td>
</tr>
<tr>
<td>2</td>
<td>21.9</td>
<td>15.6</td>
<td>30.6</td>
<td>24.6</td>
<td>24.9</td>
<td>29.1</td>
<td>32.5</td>
</tr>
<tr>
<td>3</td>
<td>1.3</td>
<td>1.3</td>
<td>1.4</td>
<td>1.6</td>
<td>1.4</td>
<td>1.2</td>
<td>1.2</td>
</tr>
<tr>
<td>4</td>
<td>32.9</td>
<td>11.9</td>
<td>24.7</td>
<td>15.2</td>
<td>20.8</td>
<td>32.1</td>
<td>25.7</td>
</tr>
<tr>
<td>5</td>
<td>0.5</td>
<td>0.6</td>
<td>0.6</td>
<td>0.8</td>
<td>0.7</td>
<td>0.6</td>
<td>0.6</td>
</tr>
<tr>
<td>6</td>
<td>35.5</td>
<td>18.5</td>
<td>23.8</td>
<td>9.3</td>
<td>19.9</td>
<td>35.7</td>
<td>32.8</td>
</tr>
<tr>
<td>7</td>
<td>2.6</td>
<td>2.3</td>
<td>2.5</td>
<td>1.9</td>
<td>2.1</td>
<td>2.5</td>
<td>2.1</td>
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<tr>
<td>8</td>
<td>39.1</td>
<td>23.4</td>
<td>26.5</td>
<td>20.1</td>
<td>21.1</td>
<td>51.4</td>
<td>44.0</td>
</tr>
<tr>
<td>9</td>
<td>0.4</td>
<td>0.4</td>
<td>0.4</td>
<td>0.5</td>
<td>0.4</td>
<td>0.2</td>
<td>0.4</td>
</tr>
<tr>
<td>10</td>
<td>20.9</td>
<td>21.1</td>
<td>18.7</td>
<td>11.7</td>
<td>17.0</td>
<td>22.6</td>
<td>20.7</td>
</tr>
<tr>
<td>11</td>
<td>0.5</td>
<td>0.8</td>
<td>0.7</td>
<td>0.7</td>
<td>0.8</td>
<td>0.5</td>
<td>0.6</td>
</tr>
<tr>
<td>12</td>
<td>31.5</td>
<td>17.2</td>
<td>23.6</td>
<td>10.6</td>
<td>15.3</td>
<td>36.6</td>
<td>32.4</td>
</tr>
<tr>
<td>13</td>
<td>0.3</td>
<td>0.4</td>
<td>0.4</td>
<td>0.5</td>
<td>0.4</td>
<td>0.3</td>
<td>0.3</td>
</tr>
<tr>
<td>14</td>
<td>35.2</td>
<td>20.5</td>
<td>17.6</td>
<td>11.0</td>
<td>17.0</td>
<td>21.3</td>
<td>37.8</td>
</tr>
<tr>
<td>15</td>
<td>1.9</td>
<td>2.3</td>
<td>1.8</td>
<td>1.6</td>
<td>1.8</td>
<td>1.9</td>
<td>1.9</td>
</tr>
<tr>
<td>16</td>
<td>22.2</td>
<td>20.4</td>
<td>28.8</td>
<td>15.1</td>
<td>20.2</td>
<td>26.3</td>
<td>23.9</td>
</tr>
</tbody>
</table>
The results of the ANOVA (Tab. 5) demonstrate that in all species, interpopulation variability is dominant in total variability.

The interpopulation variation was evaluated in *Dysphania botrys*, *D. pumilio*, *D. ambrosioides* and *D. multifida*, as *D. schraderiana* was investigated in a single population.

The values of Euclidean distances (Ed) varied within different limits in the individual taxa (Tabs. 6–9)

The UPGA cluster analysis based on the morphological pairwise similarities (Euclidean distances between population centroids) enabled us to detect two main clusters A and B.
Cluster A consists of two subclusters A1 and A2. The populations of *Dysphania multifida* form the subcluster A1, while the populations of *D. botrys*, *D. pumilio* and *D. schraderiana* were grouped by their own species in the subcluster A2. Cluster B contains the populations of *D. ambrosioides*.

Tab. 5. Percentage of the interpopulation variation in the overall morphological variation for each character in *Dysphania*

<table>
<thead>
<tr>
<th>Character No</th>
<th>Species</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td><em>D. botrys</em></td>
</tr>
<tr>
<td>1</td>
<td>37.7</td>
</tr>
<tr>
<td>2</td>
<td>29.2</td>
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<tr>
<td>3</td>
<td>65.1</td>
</tr>
<tr>
<td>4</td>
<td>46.4</td>
</tr>
<tr>
<td>5</td>
<td>69.1</td>
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<tr>
<td>6</td>
<td>61.8</td>
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<tr>
<td>7</td>
<td>66.5</td>
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<tr>
<td>8</td>
<td>67.7</td>
</tr>
<tr>
<td>9</td>
<td>45.1</td>
</tr>
<tr>
<td>10</td>
<td>76.6</td>
</tr>
<tr>
<td>11</td>
<td>69.3</td>
</tr>
<tr>
<td>12</td>
<td>52.1</td>
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<tr>
<td>13</td>
<td>66.5</td>
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<tr>
<td>14</td>
<td>55.7</td>
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<td>15</td>
<td>78.9</td>
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<td>16</td>
<td>33.1</td>
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<td>17</td>
<td>41.6</td>
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<td>18</td>
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<tr>
<td>19</td>
<td>57.0</td>
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<tr>
<td>20</td>
<td>76.8</td>
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<tr>
<td>21</td>
<td>52.5</td>
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<tr>
<td>22</td>
<td>66.4</td>
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<tr>
<td>23</td>
<td>64.0</td>
</tr>
<tr>
<td>24</td>
<td>61.7</td>
</tr>
</tbody>
</table>

Tab. 6. Values of Euclidean distances between the pairs of populations of *D. pumilio* based on 24 characters.

<table>
<thead>
<tr>
<th>Number of population</th>
<th>214</th>
<th>217</th>
</tr>
</thead>
<tbody>
<tr>
<td>217</td>
<td>54</td>
<td></td>
</tr>
<tr>
<td>221</td>
<td>111</td>
<td>57</td>
</tr>
</tbody>
</table>

Tab. 7. Values of Euclidean distances between pairs of populations of *D. ambrosioides* based on 24 characters.

<table>
<thead>
<tr>
<th>Number of population</th>
<th>32</th>
<th>104</th>
</tr>
</thead>
<tbody>
<tr>
<td>104</td>
<td>125</td>
<td></td>
</tr>
<tr>
<td>201</td>
<td>295</td>
<td>194</td>
</tr>
</tbody>
</table>

(Fig. 1). Cluster A consists of two subclusters A1 and A2. The populations of *Dysphania multifida* form the subcluster A1, while the populations of *D. botrys*, *D. pumilio* and *D. schraderiana* were grouped by their own species in the subcluster A2. Cluster B contains the populations of *D. ambrosioides*.
The results from the SDA are presented in table 10. The first three discriminant (canonical) functions account for over 98% of the total variance. Overall, the discrimination between species is highly significant (Wilks’ Lambda = 0.00026, F = 567.01, p < 0.001). Wilks’ Lambda is the standard statistic that has been used to denote the statistical significance of the discriminatory power of the current model. The closer to 0 the Wilks’ Lambda, the greater is the contribution of the model to the overall discrimination. The Partial Wilks’ Lambda values in the first column (Tab. 10) present the unique contribution of each character in the model to the discrimination between species. The lower the value in this column, the greater is the unique discriminatory power of the respective character. All discriminant functions and variables in the model were highly significant (p < 0.001)

The standardized discriminant function coefficients determine the unique contribution of each character to the discriminant functions D1–D3. The first discriminant function which provides the most overall discrimination between the five species is loaded most heavily by the width and the length of perianth lobes in the female flower, and the length of perianth lobes in the bisexual flower. The second discriminant function seems to be marked

---

**Tab. 8.** Values of Euclidean distances between the pairs of populations within *Dysphania multifida* based on 24 characters

<table>
<thead>
<tr>
<th>Number of population</th>
<th>39</th>
<th>40</th>
<th>105</th>
<th>106</th>
<th>107</th>
<th>108</th>
</tr>
</thead>
<tbody>
<tr>
<td>40</td>
<td>537</td>
<td></td>
<td></td>
<td></td>
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<td></td>
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<td>105</td>
<td>254</td>
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<td></td>
</tr>
<tr>
<td>106</td>
<td>465</td>
<td>122</td>
<td>216</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>107</td>
<td>418</td>
<td>89</td>
<td>173</td>
<td>51</td>
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<tr>
<td>108</td>
<td>40</td>
<td>497</td>
<td>215</td>
<td>425</td>
<td>378</td>
<td></td>
</tr>
<tr>
<td>109</td>
<td>158</td>
<td>388</td>
<td>96</td>
<td>310</td>
<td>266</td>
<td>121</td>
</tr>
</tbody>
</table>

**Tab. 9.** Values of Euclidean distances between the pairs of populations of *Dysphania botrys* based on 24 characters

<table>
<thead>
<tr>
<th>Number of Population</th>
<th>29</th>
<th>97</th>
<th>100</th>
<th>35</th>
<th>102</th>
<th>99</th>
<th>101</th>
<th>53</th>
<th>30</th>
<th>34</th>
</tr>
</thead>
<tbody>
<tr>
<td>97</td>
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<tr>
<td>100</td>
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<td></td>
<td></td>
</tr>
<tr>
<td>102</td>
<td>74</td>
<td>5</td>
<td>37</td>
<td>60</td>
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<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>99</td>
<td>77</td>
<td>126</td>
<td>97</td>
<td>83</td>
<td>129</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>98</td>
<td>171</td>
<td>152</td>
<td>162</td>
<td>171</td>
<td>150</td>
<td>245</td>
<td></td>
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<td></td>
</tr>
<tr>
<td>101</td>
<td>296</td>
<td>265</td>
<td>273</td>
<td>283</td>
<td>266</td>
<td>267</td>
<td>410</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>53</td>
<td>90</td>
<td>119</td>
<td>106</td>
<td>100</td>
<td>119</td>
<td>158</td>
<td>105</td>
<td>378</td>
<td></td>
<td></td>
</tr>
<tr>
<td>30</td>
<td>13</td>
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<td>15</td>
<td>63</td>
<td>84</td>
<td>166</td>
<td>290</td>
<td>92</td>
<td></td>
</tr>
<tr>
<td>34</td>
<td>151</td>
<td>152</td>
<td>153</td>
<td>155</td>
<td>151</td>
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<td>69</td>
<td>148</td>
</tr>
<tr>
<td>45</td>
<td>205</td>
<td>274</td>
<td>241</td>
<td>220</td>
<td>276</td>
<td>160</td>
<td>347</td>
<td>398</td>
<td>243</td>
<td>216</td>
</tr>
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<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

The results from the SDA are presented in table 10. The first three discriminant (canonical) functions account for over 98% of the total variance. Overall, the discrimination between species is highly significant (Wilks’ Lambda = 0.00026, F = 567.01, p < 0.001). Wilks’ Lambda is the standard statistic that has been used to denote the statistical significance of the discriminatory power of the current model. The closer to 0 the Wilks’ Lambda, the greater is the contribution of the model to the overall discrimination. The Partial Wilks’ Lambda values in the first column (Tab. 10) present the unique contribution of each character in the model to the discrimination between species. The lower the value in this column, the greater is the unique discriminatory power of the respective character. All discriminant functions and variables in the model were highly significant (p < 0.001)
GROZEVA N. H., CVETANOVA Y. G.

Fig. 1. Dendrogram of the cluster analysis of the genus *Dysphania*: No. 105–40 = *D. multifida* (m); No. 217–221 = *D. pumilio* (p); No. 97–101 = *D. botrys* (b); No. 103 = *D. schraderiana* (s); No. 201–104 = *D. ambrosioides* (a)

Tab. 10. The sequence of the ten most significant characters for discrimination between five species of genus *Dysphania* obtained by SDA and corresponding values of Partial Wilks’ Lambda, F-values and standardized coefficients for discriminant functions D₁–D₃. The eigenvalues, cumulative percentage of the variance explained by the first three roots are presented in the bottom two rows.

<table>
<thead>
<tr>
<th>Characters</th>
<th>F-remove</th>
<th>Wilks’ Lambda</th>
<th>D₁</th>
<th>D₂</th>
<th>D₃</th>
</tr>
</thead>
<tbody>
<tr>
<td>x₁₆. Width of perianth lobes in female flower</td>
<td>307.0124</td>
<td>0.000674</td>
<td><strong>–0.75486</strong></td>
<td>–0.096454</td>
<td>–0.255158</td>
</tr>
<tr>
<td>x₇. Width of upper leaf</td>
<td>169.4663</td>
<td>0.000488</td>
<td>–0.03937</td>
<td>–0.384323</td>
<td><strong>0.715588</strong></td>
</tr>
<tr>
<td>x₈. Length of upper leaf</td>
<td>137.8743</td>
<td>0.000445</td>
<td>–0.12250</td>
<td>–0.666162</td>
<td>–0.372245</td>
</tr>
<tr>
<td>x₁₅. Length of perianth lobes in female flower</td>
<td>137.6188</td>
<td>0.000445</td>
<td><strong>–0.71658</strong></td>
<td>0.390183</td>
<td>0.069843</td>
</tr>
<tr>
<td>x₁₃. Length of perianth lobes in bisexual flower</td>
<td>72.0499</td>
<td>0.000356</td>
<td><strong>0.47251</strong></td>
<td>–0.214634</td>
<td>–0.386907</td>
</tr>
<tr>
<td>x₂₄. Thickness of fruit</td>
<td>54.0739</td>
<td>0.000332</td>
<td>–0.15462</td>
<td>–0.220721</td>
<td>0.206799</td>
</tr>
<tr>
<td>x₁₁. Length of flower petiole</td>
<td>85.6233</td>
<td>0.000374</td>
<td>–0.01660</td>
<td>–0.107996</td>
<td><strong>0.471146</strong></td>
</tr>
<tr>
<td>x₄. Length of basal leaf petiole</td>
<td>62.8143</td>
<td>0.000343</td>
<td>0.10170</td>
<td>0.156215</td>
<td><strong>0.544039</strong></td>
</tr>
<tr>
<td>x₁₂. Diameter of flower</td>
<td>45.4535</td>
<td>0.000320</td>
<td>0.28564</td>
<td>–0.306588</td>
<td>–0.234661</td>
</tr>
<tr>
<td>x₉. Length of upper leaf petiole</td>
<td>37.9717</td>
<td>0.000310</td>
<td>0.10216</td>
<td>–0.273894</td>
<td>–0.373920</td>
</tr>
<tr>
<td>Eigenvalues</td>
<td>41.98092</td>
<td>7.680871</td>
<td>4.657472</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Cumulative Prop. of Variance</td>
<td><strong>0.76118</strong></td>
<td>0.900440</td>
<td><strong>0.984887</strong></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>
mostly by the length of the upper leaf, while the width of the upper leaf, length of flower petiole and length of basal leaf petiole contribute mostly to the third discriminant function.

The populations of *D. multifida*, *D. ambrosioides* and *D. schraderiana* have been classified 100% correctly on the fourth step of the SDA while the populations of the other two species *D. botrys* and *D. pumilio* were classified with 98.46% and 92.22% correctness, respectively, on the tenth step.

The scatter plot of the canonical scores of the individuals from the five species on the first two discriminant dimensions (Fig. 2) enabled us to determine the nature of the discrimination for the first two discriminant roots. The first discriminant function mostly discriminates between *D. multifida* and the four other species. Clearly, the populations of *D. multifida* are plotted much further to the right of the central line. The second function seems to provide some discrimination between the species of *D. ambrosioides*, *D. schraderiana* and *D. multifida* (which mostly show negative values for the second canonical function) and the other two species *D. botrys* and *D. pumilio* (which have mostly positive values).

Discussion

The results of the present complete population research of genus *Dysphania* confirm the data from our previous research (GROZEV Aand CVETANOV A2008, 2011) on the species from family Chenopodiaceae, namely that vegetative traits are more variable than generative ones (Tabs. 2–4). The lamina size of the lower leaves, petiole length of upper or lower leaves and stem height are the most variable traits in almost all populations. The characteristics of the seed dimensions (i.e., length, width and ratio between them) are some of the most con-
servative traits. No variability has been recorded for the trait length of female flower
perianth in two of the populations of *D. botrys*, No. 45 from Tselina and No. 99 from Tro-
yan. The high variability of stem height and leaf characters of the species from *Cheno-
podium* senso lato, and the lack of any considerable variability in seed characters have been
acknowledged by researchers of the genus from various parts of the world (KOWAL 1953,
ENGSTRAND and GUSTAFSSON 1972–1974, REYNOLDS and CRAUFORD 1980, BASSET and

The main source of phenotype variation in all species is the interpopulation variation.
The interpopulation differences were fewer and varied within much narrower limits in *D.
pumilio* (Tab. 6). That is associated with the small differences between population habitats
(Tab. 1) as well as with the lack of karyotype variability. Low variability level has been
found among the populations of *D. ambrosioides* (Tab. 7), which relates to their similar hab-
itats (sandy terrains in close proximity to a water basin) and similar altitudes (28–164 m).
The recorded higher variability for the population from the Danubian plain than in the two
populations from the Thracian lowland could be explained by either the differences in the
soil and climatic conditions due to their geographic remoteness or the established karyotype
differences.

Large interpopulation morphological variation was observed in *D. multifida* (Tab. 8)
and the greatest differences have been recorded among populations with different karyo-
types. Among populations with the same structure of the karyotypes, strong morphological
similarity was recorded between populations with similar community components – No. 39
from the island of Golyama Barzina and No. 108 from Parvomay (Ed = 40), where *D.
multifida* dominates and cereal species are quite limited; No. 106 from Ihtiman and No. 107
from Ivaylovgrad (Ed = 51), where cereal species are dominant. According to our observa-
tions, the established relationships is directly linked to the biological characteristics of the
species. *D. multifida* is a perennial grass plant with a shallow root system and highly
branched creeping stems. Its normal vegetative development needs space and efficient
amount of water and food supplies in the topsoil, and these depend on the composition of the
population and the existing relationships within it.

The largest interpopulation morphological variation was observed in *Dysphania botrys*
(Tab. 9) which is in agreement with the diversity of ecological conditions, altitudes (0–748
m), ecosystems (Tab. 1), as well as the constituted karyotype differences. The interpopulation
differences within this species were the least between populations No. 97 from Po-
morie, No. 102 from »Sinite kamani« Natural Park (Ed = 5), and population No. 29 from the
Varna lake and population No. 30 from Plovdiv (Ed = 13), which have identical karyotype
structures. High morphological similarity was found between population No. 35 from
Belene and population No. 30 from Plovdiv (Ed = 15), and population No. 29 from Varna
Lake (Ed = 22), which are with different karyotypes, although they have similar ecological
conditions (altitudes up to 200 m, flat terrain, close to a water basin with moist and sandy
soil). Of all studied populations, the greatest differences are those from Tselina (population
No 45) and Blagoevgrad (population No 101) where the species is a weed in a cultivated
community. This is related to better soil conditions (i.e., food substances, moisture, and aeri-
ation), and weaker competition between the cultivated and weed plant than in the species’
natural habitat. Regardless of the similar composition of communities, both populations dif-
fer significantly from each other, which relates to their appurtenance to two different and
quite distant floristic regions and to the differences found in their karyotypes.

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The interpopulation variability is dominant within the total variability in all species (Tab. 5), which is due mainly to differences in ecological conditions and especially in the various plant communities to which they belong, as for the populations of *D. ambrosioides*, *D. botrys* and *D. multifida* karyological variability is also a factor. The less pronounced intrapopulation variability relates to the smaller number and area of the population, and the uniform ecological conditions within its borders.

After comparing all populations of the species in the genus according to the entire morphological set of quantitative traits (Fig. 1), the greatest similarity was found between the two diploid species *D. botrys* and *D. pumilio*. Quite similar to these two is the third diploid species – *D. schraderiana*. The three species also have similar ecological preferences. A certain morphological similarity was registered between the tetraploid *D. multifida* and the three diploid species, as well as between the tetraploid *D. ambrosioides* and all other species in the genus. Such morphological similarity among the five species was also noted by UOTILA (2001) in Flora Nordica.

In spite of the high morphological similarity among the species, each one of them could be recognized correctly by certain characteristics (Fig. 2). The top ten significant features for differentiation of species, found by SDA were: width of perianth lobes in female flower, width of upper leaf, length of upper leaf, length of perianth lobes in female flower, length of perianth lobes in bisexual flower, thickness of fruit, length of flower petiole, length of basal leaf petiole, diameter of flower, length of upper leaf petiole (Tab. 10). The distinction between *D. multifida*, *D. ambrosioides* and *D. schraderiana* is based on differences in the quantitative traits, while such discrimination between *D. botrys* and *D. pumilio* is not always feasible. Due to the similarity in the greater part of the qualitative traits of these two species, qualitative characteristics should be used as well for their clear-cut distinction. *D. pumilio* could be distinguished easily from *D. botrys* as well as from the other species of the genus by its 4–5 pale green perianth leaves that harden and whiten at the fruit, as well as by reddish-brown to brown-black seeds with an imperceptible blunt edge. *D. botrys* was characterized by 5 yellowish-green perianth leaves that do not fully cover the fruit and do not harden and greyish-black seeds with an edge longitudinally split up in two. *D. schraderiana* differed by 4–5 perianth lobes with clearly defined serrated dorsal edge and black seeds with blunt edge. *D. ambrosioides* is easily recognizable by its five pale green perianth lobes with clearly defined non-serrated dorsal edge, connate up to 1/2 of their length and brown seeds with blunt edge. *D. multifida* differed to the greatest degree from the other species with its five heavily furrowed perianth lobes, with no dorsal edge, connate along their entire length, that are preserved at the fruit and form a false pod and dark brown round seeds. Each species could also be identified by its typical indentation of the leaf lamina, however, that is not always a sure method since it is necessary to collect a greater number of whole plants in order to trace the lamina changeability of the lower, middle, and upper leaves.

The data from the present study confirm that polyploidy and diploidy are basic evolutionary mechanisms in genus *Dysphania*. In tracing the morphological variability, the tendency, known for *Chenopodium* senso lato, to reduce the sizes of the generative organs and the number of perianth lobes was established. The reduced size of perianth lobes and the decrease in their number facilitate anemophilous pollination. The retention of the perianth lobes at the fruit and the presence of glands on their surface is an adaptation to exozoochory. The nuts stick to animal fur and are carried as they move around, while the reduced size of
the seeds favours their transportation at greater distances. Perianth lobes in some species have additional features securing reliable attachment: rough surface in *D. multifida*, pointed dorsal edge in *D. ambrosioides*, serrated dorsal edge in *D. schraderiana*. The lack of a dorsal edge on the perianth lobes of *D. botrys* and *D. pumilio* is perfectly compensated for by there being more glands on the entire surface of the first species, and on the upper parts of the second.

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