

## MORPHOLOGICAL, GENETICAL AND ECOGEOGRAPHICAL CHARACTERIZATION OF LONG-WINGED SPECIES OF ONOBRYCHIS SECT. ONOBRYCHIS (FABACEAE) IN IRAN

Z. Toluei, M. Atri, M. Ranjbar & M. Wink

Received 30.11.2011. Accepted for publication 08.03.2012.

Toluei, Z., Atri, M., Ranjbar, M. & Wink, M. 2012 06 30: Morphological, genetical and ecogeographical characterization of long-winged species of *Onobrychis* sect. *Onobrychis* (Fabaceae) in Iran. -Iran. J. Bot. 18 (1): 31-41. Tehran.

In this study, 30 specimens (6 populations) from long-winged species of *Onobrychis* sect. *Onobrychis* including *O. verae* Širj., *O. gontscharovii* Vassilcz., *O. ptychophylla* Širj. & Rech. f., *O. sosnovskyi* Grossh. and *O. araxina* Schischkin were collected from their natural habitats in Iran. The specimens were biometrically assessed using 45 quantitative and 15 qualitative morphological characters. In order to study the genetic variations of different species, nucleotide sequence data from the internal transcribed spacer of the nuclear rRNA genes (ITS) were obtained. Data regarding longitude, latitude, altitude, slope inclination, slope direction, substrate, min/max annual temperatures, number of rainy days, annual precipitation as well as soil traits including texture, electrical conductivity, organic carbon, total nitrogen, available phosphorus and potassium, total neutralizing value, pH, and saturation percentage were recorded in each sample site. Cluster analysis of morphological characters showed two major groups separating northwest species from northeast ones. Furthermore, the ITS data analysis results showing two main groups, which are exactly in concordance with morphological groups. Also canonical correspondence analysis (CCA) of ecological data showed the precipitation, rainy days, sand% and silt% as the most effective factors for this separation. Furthermore it showed the effect of ecological factors on each species. *Onobrychis araxina* Schischkin is recorded for the first time from Azerbaijan Gharbi province, Iran. An identification key to the long-winged species of section *Onobrychis* in Iran is presented.

Zeinab Toluei (correspondence, <[ztoluei@basu.ac.ir](mailto:ztoluei@basu.ac.ir), [ztoluei@yahoo.com](mailto:ztoluei@yahoo.com)>), Morteza Atri and Massoud Ranjbar, Department of Biology, Faculty of Sciences, Bu-Ali-Sina University, P. O. Box 65175/4161, Hamedan, Iran. - Michael Wink, Institute of Pharmacy and Molecular Biotechnology (IPMB), Dept. of Biology, Ruprecht-Karls-Universität Heidelberg, Im Neuenheimer Feld 364, 69120 Heidelberg, Germany.

**Keywords.** Ecogeography, Iran, ITS, morphology, *Onobrychis* sect. *Onobrychis*.

ویژگی‌های مورفولوژیکی، ژنتیکی و اکوجغرافیایی گونه‌های بال بلند بخش *Onobrychis* از جنس *Onobrychis* Mill. در ایران

زینب طلوعی، دانشجوی دکتری گروه زیست‌شناسی دانشگاه بوعلی سینا.

مرتضی عطری، استاد گروه زیست‌شناسی دانشگاه بوعلی سینا.

مسعود رنجبر، دانشیار گروه زیست‌شناسی دانشگاه بوعلی سینا.

مایکل وینک، استاد گروه زیست‌شناسی مؤسسه فارماکولوژی و بیوتکنولوژی مولکولی دانشگاه هایدلبرگ آلمان.

در این بررسی، ۳۰ نمونه (۶ جمعیت) از گونه‌های بال بلند بخش *Onobrychis* از جنس *Onobrychis* Mill. شامل گونه‌های *O. verae* Širj., *O. gontscharovii* Vassilcz., *O. ptychophylla* Širj. & Rech. f., *O. sosnovskyi* Grossh. و *O. araxina* Schischkin، از زیستگاه‌های طبیعی خود در ایران جمع‌آوری شدند. این نمونه‌ها با استفاده از ۴۵ ویژگی کمی و ۱۵ ویژگی کیفی مورفولوژیکی مورد ارزیابی قرار گرفتند. به منظور بررسی تنوع ژنتیکی گونه‌های مورد نظر، از داده‌های مربوط به توالی نوکلئوتیدی ناحیه ITS ژن nrRNA استفاده شد. در هر محل جمع‌آوری، اطلاعاتی شامل طول و عرض جغرافیایی، ارتفاع، جهت و میزان شیب، بستر، حداقل و حداکثر درجه حرارت سالانه، تعداد روزهای بارانی، بارش سالانه و همچنین ویژگی‌های مربوط به خاک از جمله بافت، رسانایی الکتریکی، کربن آلی، نیتروژن کل، فسفر و پتاسیم در دسترس، ارزش مواد خثی شونده، pH و درصد اشباع ثبت شد. بررسی و مقایسه نتایج حاصل از تجزیه و تحلیل خوشه‌ای داده‌های مورفولوژیکی، منجر به تشخیص دو گروه اصلی گردید. این گروه‌بندی مبین جدایی گونه‌های شمال غربی از شمال شرقی کشور

ایران است. علاوه بر این، نتایج حاصل از تجزیه و تحلیل داده‌های مربوط به توالی نوکلئوتیدی ITS نیز نتایج به دست آمده از آنالیز داده‌های مورفولوژیکی را تایید می‌کند. همچنین نتایج حاصل از آنالیز داده‌های اکولوژیکی به روش CCA، تعداد روزهای بارانی، بارش سالانه، درصد شن و درصد سیلت خاک را به عنوان مهمترین عوامل مؤثر در جدایی این دو گروه معرفی کرد. همچنین چگونگی اثر عوامل زیست محیطی را برای هر یک از گونه‌ها نشان داد. در این بررسی *O. araxina* Schischkin برای اولین بار از ایران گزارش می‌شود. کلید شناسایی گونه‌های بال بلند بخش *Onobrychis* در ایران نیز ارائه شده است.

## INTRODUCTION

The genus *Onobrychis* Miller belongs to *Fabaceae-Hedysareae* (Polhill 1981; Lock 2005). Many confusions and contradictions are found in the taxonomy of *Onobrychis*. It comprises about 130 species and is mainly distributed in the north temperate regions, but its biodiversity centers are in the eastern Mediterranean area and western Asia especially Iran and Anatolia (Lock 2005, Yildiz & al. 1999). The genus was subdivided into two subgenera and nine sections (Rechinger 1984). The section *Onobrychis* (subgenus *Onobrychis*), is the second largest section of genus *Onobrychis* in Iran with 15 species. (Rechinger 1984). The species of this section are important perennial species used for high-protein fodder for ruminants and equines, for increasing the nutritive value of drought-resistant pastures due to nitrogen fixation, and for soil conservation (Abou-El-Enain 2002; Elena 2006). This section seems to be the most problematic and the boundary of species is not completely clear due to the polymorphism in morphological characters.

In conjunction with other taxonomic studies on *Onobrychis* spp. in Iran (e.g., Ranjbar & al. 2004, 2007, 2009, 2010; Ahangarian & al. 2007; Toluei & al. 2010; Amirabadi-zadeh, 2011), the present investigations were carried out to study the interspecific relationships between different long-winged species of section *Onobrychis* including *O. verae* Širj., *O. gontscharovii* Vassilez., *O. ptychophylla* Širj. & Rech. f., *O. sosnovskyi* Grossh. and *O. araxina* Schischkin based on evidence from morphological characters and nucleotide sequence data from the internal transcribed spacer of the nuclear rRNA genes (ITS) also for showing the ecological characterization of each species. In this study, the genetic variations were investigated by means of nucleotide sequence data from the ITS region. Among the nuclear markers, ITS regions have been especially useful in recent phylogenetic studies at lower taxonomic levels in many angiosperm groups, including Fabaceae (Wojciechowski & al. 1993; Käss & Wink 1997a, b; Ainouche & Bayer 1999; Davis & al. 2002; Ahangarian & al. 2007). Ecogeographic data regarding the habitats of these species in Iran are reported here for the first time. In addition, *O. araxina* Schischkin is recorded for the first time from Iran.

Specifically, four main aims are addressed in this paper: (i) to detect the distribution of long-winged species of section *Onobrychis* in Iran; (ii) to study the morphological variation and diagnostic characters of long-winged species (iii); to assess the genetic variation of different long-winged species and (iv) to establish the relationships between species distribution and environmental gradients defined by several ecogeographic variables.

## MATERIALS AND METHODS

### Sample collection

We extensively studied the distribution of long-winged species of section *Onobrychis* in Iran: 30 specimens (6 populations) from *O. verae*, *O. gontscharovii*, *O. ptychophylla*, *O. sosnovskyi* and *O. araxina* were collected from their natural habitats (6 localities) during 2008 and 2009 (April–July). Voucher specimens are deposited in the herbarium of Bu-Ali-Sina University (BASU). The according collection sites are listed in Table 1. Identification of the specimens was performed using the regional Flora (Rechinger 1984; Hedge 1970; Boissier 1872; Grossheim 1972) and a relevant monograph (Širjaev 1926).

### Morphological data

The studied specimens (five individuals: a, b, c, d and e, in each population) were assessed by biometric study. Forty-five quantitative and fifteen qualitative morphological characters were chosen and evaluated (Table 2). For statistical analysis, the qualitative characters were initially encoded according to the multi-state method, and the related means were considered for quantitative characters and these were standardized. Phenetic analysis was carried out using MVSP Vers. 3.2 (Kovach 1985–2002) with unweighted pair-group method using arithmetic averages (UPGMA) (Sneath & Sokal 1973) and phenogram of these populations were prepared by analyzing the morphological character variations in all individuals of each population.

### Molecular data

Sequence data from nuclear ribosomal DNA internal transcribed spacers (nrDNA ITS) were obtained in order to analyze the genetic variation and species

Table 1. Localities and voucher numbers of *Onobrychis* materials studied.

Species	Locality	Voucher number	Abbreviation	Gen Bank accession numbers of ITS
<i>O. sosnovskyi</i>	Azerbaijan sharghi, Sufian to Marand	Toluei&Ranjbar 23156 (BASU)	sos	JQ780469
<i>O. araxina</i>	Azerbaijan Gharbi, before Chaldoran, Alimardan village	Toluei&Ranjbar 23157 (BASU)	ara	JQ780470
<i>O. gontsharovii</i>	Khorasan Razavi, Kalat to Mashhad	Toluei&Ranjbar 23119 (BASU)	gon	JQ780471
<i>O. ptychophylla</i>	Khorasan Razavi, Quchan, 15 km to Bajgiran	Toluei&Ranjbar 23152 (BASU)	pty	JQ780472
<i>O. verae</i>	North Khorasan, Shirvan, Kouseh bifurcate	Toluei&Ranjbar 23164 (BASU)	ver	JQ780474
<i>O. verae</i>	Khorasan Razavi, 60 km from Quchan to Dargaz, Tandooreh National Park	Toluei&Ranjbar 23165 (BASU)	ver	JQ780473

differentiation. Total genomic DNAs were isolated from dried leaf material using the standard CTAB (hexadecyl trimethyl ammonium bromide) method (Doyle & Doyle 1987). The complete nrDNA ITS region was amplified using the ITS4 and ITS5 primers (White et al. 1990). PCR amplifications were performed in 50 µl reaction volumes containing: 0.5–1 µg DNA, 10 pmol of each primer, 1.5 µl dNTPs (10 mM), 5 µl Taq polymerase buffer, 10 mg/ml bovine serum albumin (BSA), 0.8 units Taq DNA polymerase (Pharmacia Biotech, Freiburg, Germany). PCR amplifications were carried out on a thermal cycler (Biomtra, Göttingen, Germany) using the following parameters: initial denaturation at 94 °C for 5 min followed by 40 cycles of denaturation at 94 °C for 1 min, annealing at 48 °C for 1 min and extension at 72 °C for 90 s and a final extension at 72 °C for 5 min. PCR products were precipitated with 4 mol/l NH<sub>4</sub>Ac and ethanol (1:1:6) followed by centrifugation for 15 min. Sequencing was performed using the DYEnamic ET Terminator Cycle Sequencing Kit (Amersham Pharmacia Biotech, Freiburg, Germany). For each sample the nrDNA ITS region was sequenced using the ITS4 primer. Sequences were analyzed using capillary electrophoresis with a Mega-Bace™ 1000 sequencer (Molecular Dynamics Inc., Amersham Pharmacia). The nrDNA ITS sequences were aligned with Bioedit Vers. 7.0.5 (Hall 1999) and adjusted manually. The aligned data matrix was reconstructed by parsimony optimality criterion using PAUP\* ver. 4.0b10 (Swofford 2002). The heuristic search option was selected using 100 replications of simple addition sequence and TBR branch-swapping with MulTrees on and steepest descent off. Support for clades were evaluated by bootstrapping using 1000 replications with the heuristic search option, simple addition sequence and TBR branch swapping. The ITS sequence of *Hedysarum wrightianum* retrieved from GenBank (GQ246059) was

included in the analyses as an outgroup. A phylogenetic tree was drawn using Tree View (Page 1996).

#### Ecogeographic data

Data obtained from each site included: altitude, longitude, latitude, slope inclination, slope orientation, substrate, min. and max. annual temperature, rainy days per year, mean annual precipitation and soil traits including texture (sand%, silt% and clay%), electrical conductivity (EC), organic carbon (OC%), total nitrogen (Total N%), available phosphorus (av. P%), available potassium (av. K%), total neutralizing value (TNV%), pH and saturation percentage (SP). Soil samples were collected from 0–30 cm depth and measured in the soil laboratory with routine methods of soil analysis. Climatic information for each collecting site was obtained from the Islamic Republic of Iran Meteorological Organization (IRIMO), see table 3. These data were converted to numerical values for canonical correspondence analysis (CCA), a technique for direct gradient analysis. Variables constraining the scores of populations are represented by arrows, the correlations of these variables to CCA axes being proportional to length, direction, and angle of arrows to axis.

## RESULTS

### New record

***Onobrychis araxina*** Schischkin, Ber. Tomsker Staats-Univ. 80: 489 (1928).

*Specimen seen.* Azerbaijan Gharbi: Chaldoran, Alimardan village, 2002 m, Toluei & Ranjbar 23157 (BASU).

*O. araxina* belongs to the sect. *Onobrychis*. *O. araxina* is characterized by the longer wings than calyx, purplish corolla, short height and fruit denticulate on crest. It was formerly known as an endemic eastern Anatolian element which is rather close to the Iranian gathering (Hedge 1973).

Table 2. List of characters and related numerical codes used in morphological studies.

No.	Characters	Numerical code
1	Plant height	cm
2	Stem length	cm
3	Length of stem hairs	mm
4	Leaf length	cm
5	Petiole length	cm
6	Number of leaflets	In no.
7	Leaflet length	mm
8	Leaflet width	mm
9	Hair length of adaxial surface of leaflet	mm
10	Hair length of abaxial surface of leaflet	mm
11	Stipule length	mm
12	Stipule width	mm
13	Length of stipule hairs	mm
14	Inflorescence length	cm
15	Peduncle length	cm
16	Length of peduncle hairs	mm
17	Number of flowers per raceme	In no.
18	Bract length	mm
19	Bract width	mm
20	Length of bract hairs	mm
21	Pedicele length	mm
22	Length of pedicele hairs	mm
23	Calyx length	mm
24	Calyx width	mm
25	Length of calyx tube	mm
26	Length of calyx teeth	mm
27	Length of calyx hairs	mm
28	Standard length	mm
29	Standard width	mm
30	Keel length	mm
31	Keel width	mm
32	Keel claw length	mm
33	Wing length	mm
34	Wing width	mm
35	Wing claw length	mm
36	Wing auricle length	mm
37	Pod length	mm
38	Pod width	mm
39	Number of crest spines	mm
40	Spine length	mm
41	Spine width	mm
42	Number of areole per pod	In no.
43	Length of pod hairs	mm
44	Seed length	mm
45	Seed width	mm

Table 2. continued.

No.	Characters	Numerical code
46	Leaflet shape	1- oblong-elliptic, linear-oblong, oblanceolate 2- oblong-elliptic, linear-oblong 3- oblong-elliptic, oblanceolate 4- oblong-elliptic, oblanceolate, orbicular 5- oblong-elliptic, linear 6- linear-oblong
47	Hair state of adaxial surface of leaflet	1- glabrous 2- glabrous and sparse 3- glabrous, sparse and loose 4- sparse and loose 5- loose
48	Hair state of abaxial surface of leaflet	1- loose 2- loose and dense
49	Standard shape	1- obovate 2- obovate and elliptic 3- orbicular
50	Standard apex	1- retuse 2- emarginated 3- truncate 4- obtuse
51	Comparison of standard and keel size	1- equal 2- equal or standard longer than keel 3- standard longer than keel 4- keel longer than standard
52	State of stem hair	1- sparse 2- sparse-loose 3- sparse, loose and dense 4- loose 5- dense
53	Flower color	1- rose-pink 2- purplish
54	State of peduncle hair	1- sparse 2- sparse and loose 3- loose and dense 4- loose 5- dense
55	State of stipule hair	1- loose and dense 2- sparse, loose and dense 3- sparse and ciliate 4- sparse, ciliate and loose 5- sparse and loose 6- glabrous, sparse, ciliate and loose 7- sparse 8- dense
56	State of bract hair	1- ciliate, sparse and loose 2- sparse and ciliate 3- loose 4- sparse and loose 5- sparse, loose and dense 6- ciliate 7- sparse 8- dense
57	Bracteole shape	1- lanceolate 2- subulate
58	State of calyx hair	1- tube is sparse and dentate is sparse and ciliate 2- tube is sparse and dentate is sparse and loose 3- tube is sparse and dentate is loose 4- both tube and dentate are loose 5- tube is sparse and dentate is dense 6- tube is sparse and loose, dentate is loose
59	Existence of black hairs	1- black hairs on calyx and pedicel 2- black hairs on pod, calyx and pedicel
60	Pod shape	1- Crest with small teeth, disk spineless 2- Crest with small spines, disk with small spinules 3- Crest with small spines, disk with small spinules or spineless 4- crest with spines and disk with hammate teeth

**Morphologic data**

*Onobrychis verae* occurs in Middle Asia, Turkmenistan and Iran. It is a perennial plant usually erect or ascending, somewhat branching at base, with 2–7 pairs of leaflets, leaflets elliptic-oblong, linear-oblong or lanceolate, standard 8–11 mm, keel 7–11 mm, wings 6–9.5 mm, pods 5–7 mm and crest 4–6 toothed, without or with very short tubercled prickles on disc. *O. gontscharovii* occurs in Central Asia. It is a perennial plant; stem woody at base, with 2–9 pairs of leaflets, leaflets elliptic or elliptic-oblong, corolla 9–10 mm long, wings 6–7 mm, pods 7–9 mm with few hammate teeth on disc and crest 4–8 toothed. *O. ptychophylla* is a perennial plant from near Ghoochan in Khorasan Province. It has stem intensely woody, with 2–11 pairs of leaflets, leaflets linear, lanceolate or elliptic-oblong, standard 8–9 mm, keel 7–9 mm, wings 6–7 mm, pods

6–7 mm long, with thin and irregular teeth on crest. *O. sosnovskiyi* occurs in Caucasus, Turkey and Iran. It is a perennial plant usually erect, branching at base, with linear or narrowly linear leaflets, corolla 9–11 mm, wings 8–9 mm and pods 5–7 mm long, denticulate at margin. *O. araxina* occurs in Turkey and Iran. It is a perennial plant with erect-ascending stems, leaves with 7–9 pairs of linear-oblong or oblong-elliptic leaflets. Corolla purplish, standard 9–11 mm, keel 9–11 mm, wing 8–9 mm and pods 5–6 mm long, denticulate on crest and with or without short hooks on the disc.

Figure 1 represents an UPGMA phenogram with average distance coefficients resulting from analyzing the morphologic characters of five individuals of each population; apparently five groups are separated and each of the species separated from the other taxa in a

Table 3. Ecogeographical variables for different studied populations of the long-winged species of *Onobrychis* sect. *Onobrychis* in Iran.

Ecogeographical factors	species					
	sos 23156	ara 23157	gon 23119	pty 23152	ver 23164	ver 23165
Altitude (m)	1626	2002	1734	1792	1697	2155
Latitude	N 38° 20. 540'	N 39° 02. 544'	N 36° 35. 879'	N 37° 39. 621'	N 37° 39. 417'	N 37° 32. 816'
Longitude	E 45° 52. 033'	E 44° 40. 613'	E 59° 56. 836'	E 57° 49. 191'	E 57° 49. 132'	E 58° 25. 909'
Slope inclination	50%	0	35%	10%	5%	80%
Slope orientation	N	0	SE	S	NW	SE
Maximum temperature (°C)	16.5	15.5	20.2	17.5	17.5	17.5
Minimum temperature (°C)	7.6	5.3	6.6	6.8	6.8	6.8
Rainy days (In no)	96.7	118.3	77.2	90.2	90.2	66
Annual precipitation (mm)	338.3	294.5	212.6	272.4	272.4	252.7
Substrate	Soil	Soil and gravel	Soil	Soil	Soil	Soil and gravel
Texture	L	SCL	L	SIL	SIL	L
Clay%	24	26	22	15	23	14
Sand%	40	49	29	31	21	39
Silt%	36	25	49	54	56	47
EC ( $\times 10^3$ mmhos/cm)	0.448	0.439	0.31	0.323	0.342	0.457
OC%	2.55	1.81	0.29	2.16	1.05	0.23
pH	7.44	7.47	7.53	7.34	7.52	7.23
av. K (ppm)	728.6	659.4	184.5	560.4	461.5	827.6
av. P (ppm)	11.4	7.2	5.2	11	10.6	31.6
Total N%	0.25	0.18	0.02	0.21	0.1	0.02
SP	35	36.2	33.1	27.1	34.3	30
TNV%	25.5	22.5	28	8.5	23	1

single group. The northeast species including *O. ptychophylla*, *O. verae* and *O. gontsharovii* are positioned adjacently and completely separated from the long-winged species in northwest of Iran, including *O. sosnovskyi* and *O. araxina*.

**Molecular data**

The aligned complete ITS region data matrix (including outgroup) consisted of 489 characters of which 445 characters are constant. Twenty-eight variable characters are parsimony-uninformative and number of parsimony-informative characters is 16. A bootstrap consensus tree was generated by analyzing the aligned sequences of nrDNA ITS region using PAUP (Fig. 2). Our study showed that there is not intrapopulation variation for nrDNA ITS sequence.

*Onobrychis sosnovskyi* and *O. araxina* are closest species with high bootstrap value of 100 and weakly allied with other species. Also the long-winged species in northeast form a well-supported clade with bootstrap support value of 90. The ITS polymorphism in different species is indicated in Table 4. In addition to nucleotide substitutions, several diagnostic indels were observed in the aligned ITS sequence data. A one bp indel (insertion and deletion) at the position 76 and a 2- bp indel at positions 94-95 occurred in the northeast species clade and deleted in *O. sosnovskyi* and *O. araxina*. Also a one base-pair (bp) indel situated at position 416 of the ITS alignment was inserted in *O.*

*sosnovskyi* and *O. araxina* but deleted in northeast species.

**Ecological data**

Ecological direct gradient analysis, as shown by CCA (Fig. 3) can establish the relative importance of each environmental variable in the distribution of the species. This figure shows the separation of the long-winged species in northeast and that ones in northwest of Iran, also precipitation, rainy days, sand% and silt% as the most effective factors for this separation. It shows that rainy days as well as the texture are the most effective factors for separation of *O. araxina*. Furthermore, it shows the precipitation being a prominent ecological factor for the separation of *O. sosnovskyi*. The slope orientation, SP and TNV are the most effective factors on *O. ptychophylla* and the most effective factor on *O. gontsharovii* is max. temperature. This figure introduces the number of rainy days, slope inclination and orientation as the most effective factor in species distribution and substrate, altitude and av. K are the lowest effective factors in species distribution. Table 5 includes interser correlations of canonical coefficients for twenty ecological variables. It should be noted that CCA axis 1 is positively correlated with OC and Total N also negatively correlated with slope inclination and orientation. CCA axis 3 is positively correlated with Total N and negatively correlated with slope orientation.

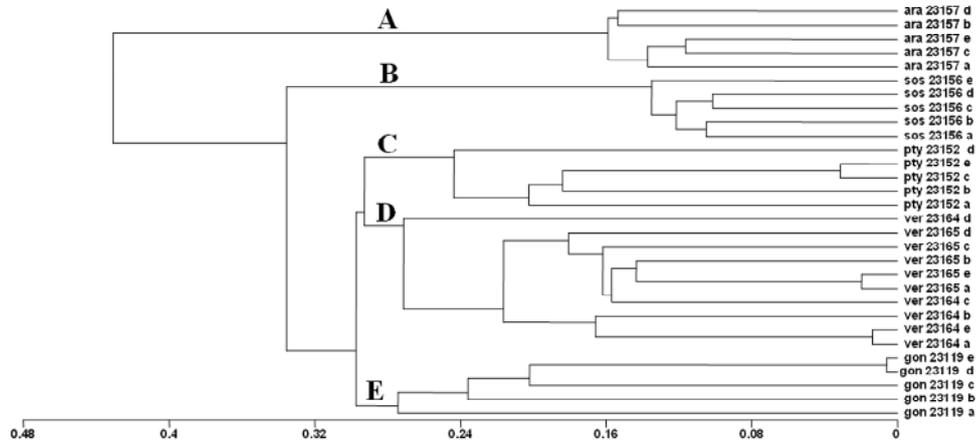


Fig. 1. The UPGMA phenogram based on the analyzing the morphological data in five individuals of each population of the long-winged species of *Onobrychis* sect. *Onobrychis* in Iran representing five groups A, B, C, D and E.

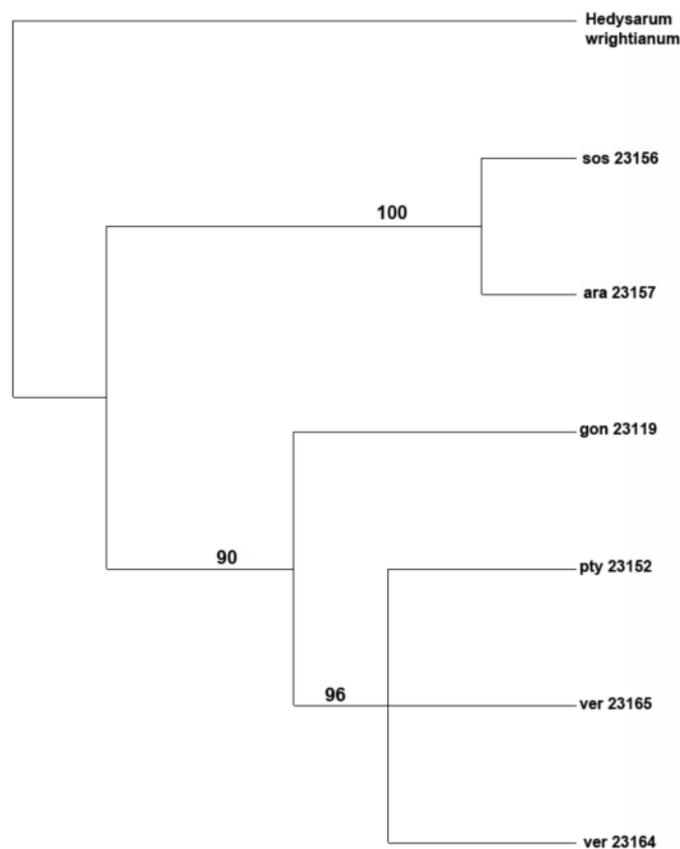


Fig. 2. A bootstrap consensus tree based on ITS sequences data of the long -winged species of *Onobrychis* sect. *Onobrychis* in Iran (abbreviations are as listed in Table 1). Numbers refer to bootstrap values. Bootstrap values greater than 50% were shown above the branches. (from 1000 replications).

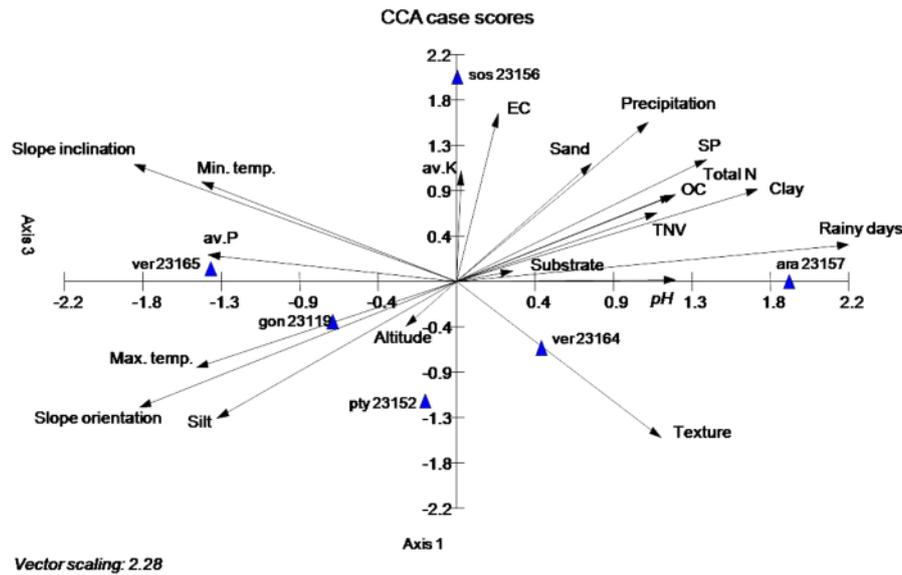


Fig. 3. Canonical correspondence analyses (CCA) of twenty ecogeographic data for collection sites of the long-winged species of *Onobrychis* sect. *Onobrychis* in Iran.

Table 4. ITS polymorphism in different populations of *Onobrychis verae*, *O. gontsharovii*, *O. ptychophylla*, *O. sosnovskyi* and *O. araxina* (abbreviations are as listed in Table 1). R= A or G, W=T or A, (-) = indel. Only sites that differed are shown.

Population	Position in alignment																								
	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	2	2	2	3	4	4	4	4	
sos 23156	A	G	T	T	C	A	C	T	G	C	C	G	-	-	-	T	G	T	C	T	G	C	C	T	A
ara 23157	A	G	T	T	C	T	C	T	G	C	C	G	-	-	-	W	G	T	C	T	G	C	C	T	A
gon 23119	G	T	C	T	C	T	C	A	C	T	T	G	G	A	T	T	G	C	T	C	T	-	T	C	G
pty 23152	G	T	C	C	T	T	G	A	C	T	C	A	G	A	T	T	G	C	C	C	T	-	T	C	G
ver 23165	G	T	C	C	C	T	G	A	C	T	C	A	G	A	T	T	G	C	C	C	T	-	T	C	G
ver 23164	G	T	C	C	C	T	G	A	C	T	C	A	G	A	T	T	R	C	C	C	T	-	T	C	G

**DISCUSSION**

In this study, cluster analysis of morphological characters in long-winged species of sect. *Onobrychis* showed two major groups separating northwest species (*O. sosnovskyi* and *O. araxina*) from northeast ones (*O. ptychophylla*, *O. verae* and *O. gontsharovii*). Based on morphological variations, each of the species separated from the other taxa (groups A, B, C, D and E in Fig. 1). The following key represents the distinguishing features of each species.

1. Plant with black hairs on pedicel, calyx and pod; calyx length 3–5 mm; wing 8–9 mm; adaxial surface of

leaflets glabrous; pods with small teeth on crest and disk spineless 2

Plant with black hairs on pedicel and calyx; calyx length 4.5–8 mm; wing 6–9.5 mm; adaxial surface of leaflets with loose and sparse hairs and rarely glabrous; pods with spines on crest and disk with small spinules, hammate teeth or spineless 3

2. Plant height 14–30 cm; flower color purplish; standard as long as keel; leaf length 1.5–8 cm

1. *O. araxina*

Plant height 35–60 cm; flower color is rose-pink; standard is longer than keel; leaf length 4–15 cm

2. *O. sosnovskyi*

Table 5. Interset correlations for twenty environmental variables.

Variables	Interset correlation		
	Axis 1	Axis 2	Axis 3
Altitude	0.015	0.017	-0.016
Slope inclination	-0.503	-0.064	0.189
Slope orientation	-0.511	-0.079	-0.297
Max. temp. (°C)	-0.001	-0.008	-0.022
Min. temp. (°C)	-0.015	-0.019	0.004
Rainy days (In no)	0.055	-0.001	-0.007
Precipitation (mm)	0.029	0.005	0.002
Substrate	0.243	1.379	0.064
Texture	0.136	-0.022	-0.149
Clay%	0.070	-0.021	0.014
Sand%	0.034	0.049	0.024
Silt%	-0.032	-0.025	-0.049
EC	0.033	0.099	0.084
OC%	0.348	-0.147	0.140
pH	0.020	0.000	-0.013
av. K	0.028	0.052	0.017
av. P	-0.094	0.132	0.019
Total N%	0.393	-0.165	0.218
SP	0.033	0.002	0.001
TNV%	0.234	-0.248	0.040

3. Pods with spines (0.3–3 mm) on crest and disk with hamate teeth; wing 6–7.5 mm; standard as long as or shorter than keel

3. *O. gontsharovii*

Pods with spines (0.2–2 mm) on crest and disk with small spinules or spineless; wing 6–9.5 mm; standard as long as or longer than keel

4

4. Plant height 50–80 cm; wing 7–9.5 mm; pods with spines (0.2–2 mm) on crest

4. *O. verae*

Plant height 15–55 cm; wing 6–7 mm; pods with spines (0.2–1 mm) on crest

5. *O. ptychophylla*

Furthermore, the ITS data analysis results showing two main groups, which are exactly in concordance with morphological groups because it separating northwest species (*O. sosnovskyi* and *O. araxina*) with high bootstrap value of 100 from northeast ones (*O. ptychophylla*, *O. verae* and *O. gontsharovii*). Also the long-winged species in northeast form a well-supported clade with bootstrap support value of 90. But these two well-supported clades (northwest and northeast ones) weakly allied with each other with bootstrap value less than 50%. Hence, it seems that the long-winged species in northeast and that ones in northwest of Iran cannot have a progenitor-derivative relationship. More studies are needed to confirm this result.

In this study, it was shown that nucleotide sequence data from the internal transcribed spacer of the nuclear rRNA genes (ITS) could be applied to investigate *Onobrychis* genetics, as indicated by other relevant studies (e.g., Käss & Wink 1997a; Ahangarian & al. 2007). It is shown to be of high potential to reveal genotypic diversity and in the longer term, to provide potential molecular markers that could be linked to

phenotypic properties. The results obtained here suggest a substantial genetic diversity across the long-winged species of sect. *Onobrychis*. Also they were found to be variable in terms of morphological characteristics.

Morphological characters are the outer, visibly expressed features of an organism – in plants these are not only affected by genetic factors, but by environmental factors as well (Mal & Doust 2005). Studies on the morphological variations according to habitat differences can help us to understand the manner, mechanism and determining factors of plant adaptation and evolution (Yang 1991). The present study shows that long-winged species of *Onobrychis* sect. *Onobrychis* exist under the following ecogeographic conditions: soil textures being loam (L), sandy clay loam (SCL) and silt loam (SIL), ranging from pH 7.23–7.53, EC 0.31–0.457, OC 0.23–2.55, clay 14–26%, sand 21–49%, silt 25–56%, av. K 184.5–827.6, av. P 5.2–31.6, Total N 0.02–0.25%, SP 27.1–36.2 and TNV 1–28%; substrate is soil or a mixture of soil and gravel; They exist on substrate without slope (0) or in N, NW, S and SE slope orientations with 5–80% inclination; lowest altitude 1626 m a.s.l., highest altitude 2155 m a.s.l.; max. annual temperature ranging from 15.5–20.2°C, min. annual temperature from 5.3–7.6°C; number of rainy days/year 66–118.3; annual precipitation 212.6–338.3 mm. CCA analysis introduced the number of rainy days, slope inclination and orientation as the main factors associated with the distribution of long-winged species of *Onobrychis* sect. *Onobrychis*. Furthermore it showed correlations

between two major morphologic-genetic groups and certain ecogeographical factors. It shows the separation of the long-winged species in northeast and that ones in northwest of Iran also precipitation, rainy days, sand% and silt% as the most effective factors for this separation. Thus there are ecogeographical patterns in the morphological and genetical variation of long-winged species of *Onobrychis* sect. *Onobrychis*.

## ACKNOWLEDGMENTS

This research was financially supported by Bu-Ali-Sina University and in part by IPMB, Heidelberg University, Germany. The authors are grateful to Dr. Javier Gonzalez for fruitful discussions and help in the molecular laboratory and to Hedwig Sauer-Gürth for extensive technical help.

## REFERENCES

- Abou-El-Enain, M. M. 2002: Chromosomal criteria and their phylogenetic implications in the genus *Onobrychis* Mill. sect. *Lophobrychis* (Leguminosae), with special reference to Egyptian species. -*Botanical Journal of the Linnean Society* 139: 409–414.
- Ahangarian, S., Kazempour Osaloo, S. & Maassoumi, A. A. 2007: Molecular phylogeny of the tribe Hedysareae with special reference to *Onobrychis* (Fabaceae) as inferred from nrDNA ITS sequences. -*Iranian Journal of Botany* 13: 64–74.
- Ainouche, A. K. & Bayer, R. J. 1999: Phylogenetic relationships in *Lupinus* (Fabaceae: Papilionoideae) based on internal transcribed spacer sequences (ITS) of nuclear ribosomal DNA. -*American Journal of Botany* 86 (4): 590–607.
- Amirabadi-zadeh, H. 2011: New records of Hedysareae (Papilionaceae) from Iran. -*Iranian Journal of Botany* 17 (1): 63–68.
- Boissier, P. E. 1872: *Onobrychis* in *Flora Orientalis* vol. 2: 525–553. -Genevae & Basileae.
- Davis, C. C., Fritsch, P. W., Li, J. & Donoghue, M. J. 2002: Phylogeny and biogeography of *Cercis* (Fabaceae): Evidence from nuclear ribosomal ITS and chloroplast *ndhF* sequence data. -*Systematic Botany* 27 (2): 289–302.
- Doyle, J. J. & Doyle, J. L. 1987: A rapid DNA isolation procedure for small quantities of fresh leaf tissue. -*Phytochemical Bulletin* 19: 11–15.
- Elena, T. 2006: Cytological aspects of the *Onobrychis* genus. -*Bulletin USAMV* 62: 154–158.
- Grossheim, A. A. 1972: *Onobrychis*. In: *Flora URSS*, eds. Shishkin, B. K. & Bobrov, E. G., vol. 13: 244–281. -Moscow, Leningrad. (English translation).
- Hall, T. A. 1999: BioEdit: a user-friendly biological sequence alignment editor and analysis program for Windows 95/98/NT. -*Nucleic Acids Symposium Series* 41: 95–98.
- Hedge, I. C. 1970: *Onobrychis* in *Flora of Turkey and the East Aegean Islands* vol. 3: 560–589. -University Press. Edinburgh.
- Käss, E. & Wink, M. 1997a: Phylogenetic relationships in the Papilionoideae (Family Leguminosae) based on nucleotide sequences of cpDNA (*rbcL*) and ncDNA (ITS 1 and ITS 2). -*Molecular Phylogenetics and Evolution* 8: 65–88.
- Käss, E. & Wink, M. 1997b: Molecular phylogeny and phylogeography of *Lupinus* (Leguminosae) inferred from nucleotide sequences of the *rbcL* gene and ITS 1 + 2 regions of rDNA. -*Plant Systematics and Evolution* 208: 139–167.
- Kovach, W. L. 1985–2002: -MVSP Vers. 3.2 3.2 ed, Kovach Computing Services. Institute of Earth Studies, University College of Wales, ABERYSTWYTH, (Shareware) Available from <http://www.kovcomp.com/MVPs/down12.html>.
- Lock, J. M. 2005: Tribe Hedysareae. In: *Legumes of the world*, eds. Lewis, G., Schrire, B., Mackinder, B. & Lock, M., vol. 489–495. -Royal Botanical Gardens, Kew. London.
- Mal, T. K. & Doust, J. L. 2005: Phenotypic plasticity in vegetative and reproductive traits in an invasive weed, *Lythrum salicaria* (Lythraceae), in response to soil moisture. -*American Journal of Botany* 92: 819–825.
- Page, R. D. M. 1996: TREEVIEW: An application to display phylogenetic trees on personal computers. -*Computer Applications in the Biosciences* 12: 357–358.
- Polhill, R. M. 1981: Hedysareae DC. In: *Advances in legume systematics I*, eds. Polhill, R. M. & Raven, P. H., vol. 367–370. -Royal Botanic Gardens, Kew. London.
- Ranjbar, M., Amirabadizadeh, H., Karamian, R. & Ghahremani, M. A. 2004: Notes on *Onobrychis* sect. *Heliobrychis* (Fabaceae) in Iran. -*Willdenowia* 34: 187–190.
- Ranjbar, M., Karamian, R., Tolui, Z. & Amirabadizadeh, H. 2007: *Onobrychis assadii* (Fabaceae), a new species from Iran. -*Annales Botanici Fennici* 44: 481–484.
- Ranjbar, M., Karamian, R. & Hadadi, A. 2009: Biosystematic study of *Onobrychis vicifolia* Scop. and *Onobrychis altissima* Grossh. (Fabaceae) in Iran. -*Iranian Journal of Botany* 15(1): 85–95.
- Ranjbar, M., Karamian, R. & Vitek, E. 2010: *Onobrychis dushanbensis* sp. nov. endemic to

- Tajikistan. -Nordic Journal of Botany 28(2): 182–185.
- Rechinger, K. H. 1984: *Onobrychis*. In: *Flora Iranica*, ed. Rechinger, K. H., no. 157: 387–464. -Akademische Druck- u. Verlagsanstalt Graz and Wien.
- Širjaev, G. 1926: *Onobrychis* generis revisio critica. -Publication de la faculte' des sciences des L'Universite Masaryk (Brno). Brno.
- Sneath, P. H. A. & Sokal, R. R. 1973: *Principles of numerical taxonomy*. -Freeman. San Francisco.
- Swofford, D. L. 2002. *PAUP\* Phylogenetic Analysis Using Parsimony (\*And Other Methods)*. Version 4.10b. Sunderland: Sinauer Associates.
- Toluei, Z., Atri, M. & Ranjbar, M. 2010: Taxonomic study of *Onobrychis transcaspica* V. Nikitin (Fabaceae) in northeastern of Iran with emphasis on altitudinal effect on morphological characters using floristic marker. -*Taxonomy and Biosystematics* 2(3): 25–38.
- White, T. J., Bruns, T., Lee, S. & Taylor, J. 1990: Amplification and direct sequencing of fungal ribosomal genes for phylogenetics. In: *PCR protocols: a guide to methods and applications*, eds. Innis, M., Gelfand, D., Sninsky, J. & White, T., vol. 315–322. -Academic Press. San Diego, California, USA.
- Wojciechowski, M. F., Sanderson, M. J., Baldwin, B. G. & Donoghue, M. J. 1993: Monophyly of aneuploid *Astragalus* (Fabaceae): evidence from nuclear ribosomal DNA internal transcribed spacer sequences. -*American Journal of Botany* 80: 711–722.
- Yang, J. 1991: Intraspecific variation in plant and the exploring methods. -*Journal of Wuhan Botanical Research* 9: 185–195.
- Yildiz, B., Ciplak, B. & Aktoklu, E. 1999: Fruit morphology of sections of the genus *Onobrychis* Miller (Fabaceae) and its phylogenetic implications. -*Israel Journal of Plant Sciences* 47: 269–282.