

KARYOMORPHOLOGICAL STUDY OF SEVEN SPECIES OF THE GENUS *ASTRAGALUS* (FABACEAE) FROM IRAN

Y. Shamsolshoara, H. Javadi & S. M. Miri

Received 2020. 01. 11; accepted for publication 2020. 07. 10

Shamsolshoara, Y., Javadi, H. & Miri S. M. 2020: Karyomorphological study of seven species of the genus *Astragalus* from Iran. -*Iran. J. Bot.* 26 (2): 172-178. Tehran.

Karyomorphological study was carried out on 11 accessions from 6 sections belonging to 7 species of *Astragalus* growing in Iran. Chromosome numbers and karyotypes of two species (*A. cephalanthus* DC. and *A. jacobsii* Podlech) are new records. Two ploidy levels $2n = 2x = 16$ and $2n = 4x = 32$ were determined, the chromosomes size varied from 2.51 in *A. cyclophyllus* to 3.79 μm in *A. fasciculifolius* Boiss. in two accessions of *A. cyclophyllus*, a single pair satellite chromosomes were observed.

Yasamin Shamsolshoara, Department of Horticulture, Karaj Branch, Islamic Azad University, Karaj, Iran. -Hamideh Javadi, Research Institute of Forests and Rangelands, Agricultural Research, Education and Extension Organization (AREEO), Tehran, Iran. - Seied Mehdi Miri (correspondence <smmiri@kiaau.ac.ir>), Department of Horticulture, Karaj Branch, Islamic Azad University, Karaj, Iran.

Key words: Chromosome number; karyotype; cytogenetic; diversity; Fabaceae

مطالعه کاربومورفولوژیکی هفت گونه از جنس گون در ایران

یاسمین شمس‌الشعرا: دانشجوی دکتری گروه باغبانی، واحد کرج، دانشگاه آزاد اسلامی، کرج، ایران

حمیده جوادی: استادیار مؤسسه تحقیقات جنگل‌ها و مراتع کشور، سازمان تحقیقات و آموزش کشاورزی، تهران، ایران

سید مهدی میری: دانشیار گروه باغبانی، واحد کرج، دانشگاه آزاد اسلامی، کرج، ایران

مطالعه سیتولوژیکی بر روی ۱۱ نمونه متعلق به ۶ بخش از ۷ گونه *Astragalus* ایران انجام شد. تعداد کروموزوم‌ها و کاریوتایپ‌های دو گونه (*A. cephalanthus* DC. و *A. jacobsii* Podlech) گزارش‌های جدیدی هستند. دو سطح پلوئیدی دی و تتراپلوئیدی $2n = 2x = 16$ و $2n = 4x = 32$ تعیین و اندازه کروموزوم‌ها از $2/51$ میکرومتر در *A. cyclophyllus* G. Beck تا $3/79$ میکرومتر در *A. ficiculifolius* Boiss. متغیر بود. در دو نمونه از گونه *A. cyclophyllus*، یک جفت ماهواره مشاهده شد.

INTRODUCTION

Astragalus L. (Fabaceae), as the largest genus of the flowering plants, contains an estimated number of 2500-3000 annual and perennial herbaceous or subshrub taxa, grouped in 150-250 sections (Lock & Simpson 1991; ILDIS 2002; Mabberley 1997; Maassoumi 1998). This genus is widely distributed throughout the temperate and arid regions of the world, and is principally located in Asia (1500 species), North America (500 species), South America (150 species) and Europe (120 species), but also on mountains in

Africa. However, the center of origin and biodiversity of *Astragalus* is Eurasia, especially the mountainous parts of South-Western and South-Central Asia (Lock & Simpson 1991; Maassoumi 1998). Iran is rich in biodiversity due to its diverse climatic and geographical conditions and with more than 820 published and valid species of *Astragalus* with an endemic rate of 57%, it is one of the most important center of diversity for this genus (Bagheri & al. 2011; Ghahremaninejad 2015; Miri & Shamsolshoara 2019; Ranjbar & al. 2014).

The first deliberate study on the chromosomes of *Astragalus* was made by Ledingham (1960) who reported the species from the old world have a basic chromosome number of $x = 8$, while those from the new world have $x = 11, 12$ and 13 (Badr and Sharawy 2007). Numerous cytological studies have been conducted on *Astragalus* species and the basic chromosome numbers of the genus has been reported as $x = 6, 7, 8, 9, 11, 15$ (Bozkurt & al. 2019). In addition, five ploidy levels ($2x, 4x, 6x, 8x$ and $12x$) are present in the genus (Ranjbar & al. 2014). The basic number of $x = 8$ is the primary basic number in *Astragalus*. It seems that $x = 7$ and $x = 6$ basic numbers have been derived from $x = 8$ by aneuploid loss of chromosomes (Badr and Sharawy 2007).

In this study, we explore the chromosome number and karyomorphology of 11 accessions belonging to 7 species of the *Astragalus* L. in Iran.

MATERIALS AND METHODS

Eleven *Astragalus* accessions belonging to 7 species from the collections of Natural Resources Gene Bank of Iran, Research Institute of Forests and Rangelands (RIFR), Tehran, Iran were included in this study (table 1). Voucher specimens have been deposited in the herbarium of Research Institute of Forests and Rangelands (TARI). The seeds were disinfected by 0.2% carboxin thiram fungicide (Vitavax®) for 3 minutes and placed in petri dishes on moist filter paper at 4°C for two weeks, then they were germinated at 25°C in a growth chamber under dark conditions. For cytological investigations, 1-1.5 cm long young root tips were pretreated with 1% α -bromonaphthalin solution for 3 h at room temperature and fixed in freshly Lewitsky solution, a mixture of 1:1 chromic acid 1%: formaldehyde 10% for 24 h at 4°C. Root tips were hydrolyzed for 15 min in 1 N HCl at 60°C, washed briefly in ddH₂O and stained in 1% hematoxylin solution for 24 h at room temperature. Squashed preparation was made in a drop of 45% acetic acid. After obtaining the appropriate metaphase, the photographs of at least 10 well-spread chromosomes were taken using digital camera mounted on an Olympus BX41 microscope and chromosomal measurements were made from five mitotic metaphases cells. Based on the measurements of chromosome length and arm ratios, a karyotype formula has been calculated as proposed by Levan & al. (1964). The symmetry class (Sc) of the karyotypes was made according to Stebbins (1971). The karyotype asymmetry was estimated using intrachromosomal (A_1) and interchromosomal (A_2) asymmetry indices suggested by Zarco (1986).

RESULTS AND DISCUSSION

The cytological data for the examined accessions is summarized in tables 2 & 3 and their karyotypes and karyograms are illustrated in fig. 1. Basic chromosome number of $x = 8$ are found in all studied accessions. The basic chromosome number reported for the old world *Astragalus* species (both from Iran and outside) is $x = 8$ and di-, tri-, tetra- and hexaploidy levels were observed (Ranjbar & Mahmoudian 2015; Sheidai & al. 2009). In our study, four species were diploid (*A. cephalanthus* DC., *A. curvirostris* Boiss., *A. dactylocarpus* Boiss. and *A. megalotropis* Bunge) and two species (*A. fasciculifolius* Boiss. and *A. jacobsii* Podlech) were tetraploid. In the accessions of *A. cyclophyllos* G. Beck, both di- and tetraploidy were observed. Therefore, both diploid and tetraploid levels occur in a single species which is also present in some other *Astragalus* species such as *A. iranicus* Bunge (Ranjbar & Mahmoudian 2015) and *A. camptoceras* Bunge (Gedik & al. 2019). These reports indicate the role of polyploidy in the evolution of *Astragalus* (Sheidai & al. 2009). As far as we know, the somatic chromosome numbers and karyotypes of two species (*A. cephalanthus* DC. and *A. jacobsii* Podlech) are reported for the first time. Chromosome numbers of the other studied accessions were consistent with previous counts (Badr & Sharawy 2007; Goldblatt & Johnson 2010; Maassoumi 1987; Marhold & Breitwieser 2009; Ranjbar & al. 2011).

The average chromosome length (CL) varies about 1.5 times between accessions (table 2). *A. cyclophyllos* (107732 TARI) exhibit much shorter chromosomes (2.51 μm) and lower total haploid chromosome length (20.71 μm), compared to other accessions. Meanwhile, longest CL (3.79 μm) and TCL (60.68 μm) have been found in *A. fasciculifolius* Boiss. (107733 TARI). Sheidai & Ghahremaninejad (2009), Sheidai & al. (2009) and Martin & al. (2019) reported the chromosome length of *Astragalus* species in the range of 1.87-5.20 μm , 2.20-5.51 μm and 2.00-3.71 μm , respectively. The chromosome length of the examined *Astragalus* accessions are generally medium size. Although the highest mean chromosome length was observed in a tetraploid accession, no significant correlation was found between mean chromosome length and polyploidy ($r = 0.05$, data not shown). However, Sheidai & al. (2009) reported that with increase in polyploidy level, decrease in the mean chromosome length has occurred (although not significant), which may indicate the loss of DNA during the polyploidy induction and species diversification.

Table 1. Geographical information of the *Astragalus* species, investigated in the present study.

Species	Section	Location	Latitude	Longitude	Altitude (m)	Herbarium code (TARI)
<i>A. cephalanthus</i> DC.	<i>Microphysa</i> Bunge	Kerman-Bardsir	N 29° 51' 00"	E 56° 05' 00"	2583	105647
<i>A. curvirostris</i> Boiss.	<i>Incani</i> DC.	Zanjan-Mahnesan	N 36° 40' 00"	E 47° 40' 00"	1950	105636
<i>A. cyclophyllos</i> G. Beck	<i>Incani</i> DC.	Zanjan-Zanjan	N 36° 40' 44"	E 48° 29' 56"	1638	107732
<i>A. cyclophyllos</i> G. Beck	<i>Incani</i> DC.	Kerman-Kerman	N 30° 17' 36"	E 57° 05' 30"	1755	107734
<i>A. cyclophyllos</i> G. Beck	<i>Incani</i> DC.	Isfahan-Fardin	N 33° 13' 30"	E 49° 58' 35"	2700	105645
<i>A. cyclophyllos</i> G. Beck	<i>Incani</i> DC.	Isfahan-Chadegan	N 32° 43' 00"	E 50° 43' 00"	2060	107735
<i>A. dactylocarpus</i> Boiss.	<i>Chronopus</i> Bunge	Yazd-Harat	N 29° 56' 00"	E 54° 05' 00"	1850	105648
<i>A. fasciculifolius</i> Boiss.	<i>Poterion</i> Bunge	Khuzestan-Haftkel	N 31° 26' 49"	E 49° 31' 46"	600	107733
<i>A. fasciculifolius</i> Boiss.	<i>Poterion</i> Bunge	Hormozgan-Bandar Abbas	N 27° 48' 15"	E 56° 17' 52"	900	105659
<i>A. jacobsii</i> Podlech	<i>Malacothrix</i> Bunge	Markazi-Tafresh	N 34° 33' 00"	E 49° 48' 00"	2050	105639
<i>A. megalotropis</i> Bunge	<i>Alopecuriodei</i> DC.	Hamedan-Malayer	N 34° 37' 05"	E 48° 43' 20"	2350	105654

Table 2. Karyotypic parameters measured on the 11 accessions of studied *Astragalus* species.

Species	2n	TCL (µm)	CL (µm)	LA (µm)	SA (µm)
<i>A. cephalanthus</i> DC. (105647)	2n=2x=16	28.03	3.50ab**	2.12ab**	1.39ab**
<i>A. curvirostris</i> Boiss. (105636)	2n=2x=16	27.38	3.42ab	2.14ab	1.29ab
<i>A. cyclophyllos</i> G. Beck (107732)	2n=2x=16	20.71	2.51c	1.52c	0.99bc
<i>A. cyclophyllos</i> G. Beck (107734)	2n=4x=32	46.73	2.92bc	1.78bc	1.14b
<i>A. cyclophyllos</i> G. Beck (105645)	2n=4x=32	58.77	3.36ab	2.10ab	1.31ab
<i>A. cyclophyllos</i> G. Beck (107735)	2n=2x=16	24.82	3.12bc	1.97b	1.14b
<i>A. dactylocarpus</i> Boiss. (105648)	2n=2x=16	21.27	2.74bc	1.86b	0.88c
<i>A. fasciculifolius</i> Boiss. (107733)	2n=4x=32	60.68	3.79a	2.33a	1.46a
<i>A. fasciculifolius</i> Boiss. (105659)	2n=4x=32	46.33	2.89bc	1.77bc	1.12b
<i>A. jacobsii</i> Podlech (105639)	2n=4x=32	43.02	2.69bc	1.67bc	1.02bc
<i>A. megalotropis</i> Bunge (105654)	2n=2x=16	28.64	3.58ab	2.11ab	1.47a

** : means within a column followed by the same letter are not significantly different according to the Duncan's multiple range test at $p \leq 0.01$.

Table 3. Asymmetrical parameters measured on the 11 accessions of studied *Astragalus* species.

Species	Sc	A ₁	A ₂	TF%	AR	CI	DRL	KF
<i>A. cephalanthus</i> DC. (105647)	2A	0.32b*	0.16	39.78b**	1.62c**	0.40ab**	5.98abc*	6m+2sm
<i>A. curvirostris</i> Boiss. (105636)	2A	0.39ab	0.15	37.27bc	1.78ab	0.37bc	5.26abc	3m+5sm
<i>A. cyclophyllos</i> G. Beck (107732)	1A	0.34b	0.14	39.49b	1.56c	0.40ab	4.39bc	5m+3sm
<i>A. cyclophyllos</i> G. Beck (107734)	2A	0.34b	0.14	39.03b	1.65c	0.39ab	3.15bc	12m+4sm
<i>A. cyclophyllos</i> G. Beck (105645)	2A	0.36ab	0.17	38.62b	1.75ab	0.38bc	4.74bc	7m+1m ^{sat} +8sm
<i>A. cyclophyllos</i> G. Beck (107735)	3A	0.39ab	0.18	36.70bc	1.82ab	0.37bc	6.21ab	3m+1m ^{sat} +4sm
<i>A. dactylocarpus</i> Boiss. (105648)	3A	0.49a	0.22	32.37c	2.16a	0.33c	6.63ab	1m+7sm
<i>A. fasciculifolius</i> Boiss. (107733)	2B	0.35b	0.23	38.50b	1.67c	0.39bc	4.56bc	9m+7sm
<i>A. fasciculifolius</i> Boiss. (105659)	2A	0.34b	0.19	38.81b	1.60c	0.39bc	4.76bc	12m+4sm
<i>A. jacobsii</i> Podlech (105639)	2A	0.38ab	0.12	37.80bc	1.69bc	0.38bc	2.33c	8m+8sm
<i>A. megalotropis</i> Bunge (105654)	1A	0.29b	0.20	41.07a	1.50c	0.41a	7.10a	7m+1sm

2n=somatic chromosome number; SC= symmetry classes of Stebbins; A₁= intra-chromosomal index; A₂= inter-chromosomal index; TF%= total form percentage; DRL=Difference of Relative Length; KF=karyotype formula; CI= Centromeric index; TL= total chromosome length; LA= long arm; SA= short arm; AR= arm ratio

* and **: means within a column followed by the same letter are not significantly different according to the Duncan's multiple range test at $p \leq 0.05$ and 0.01 , respectively.

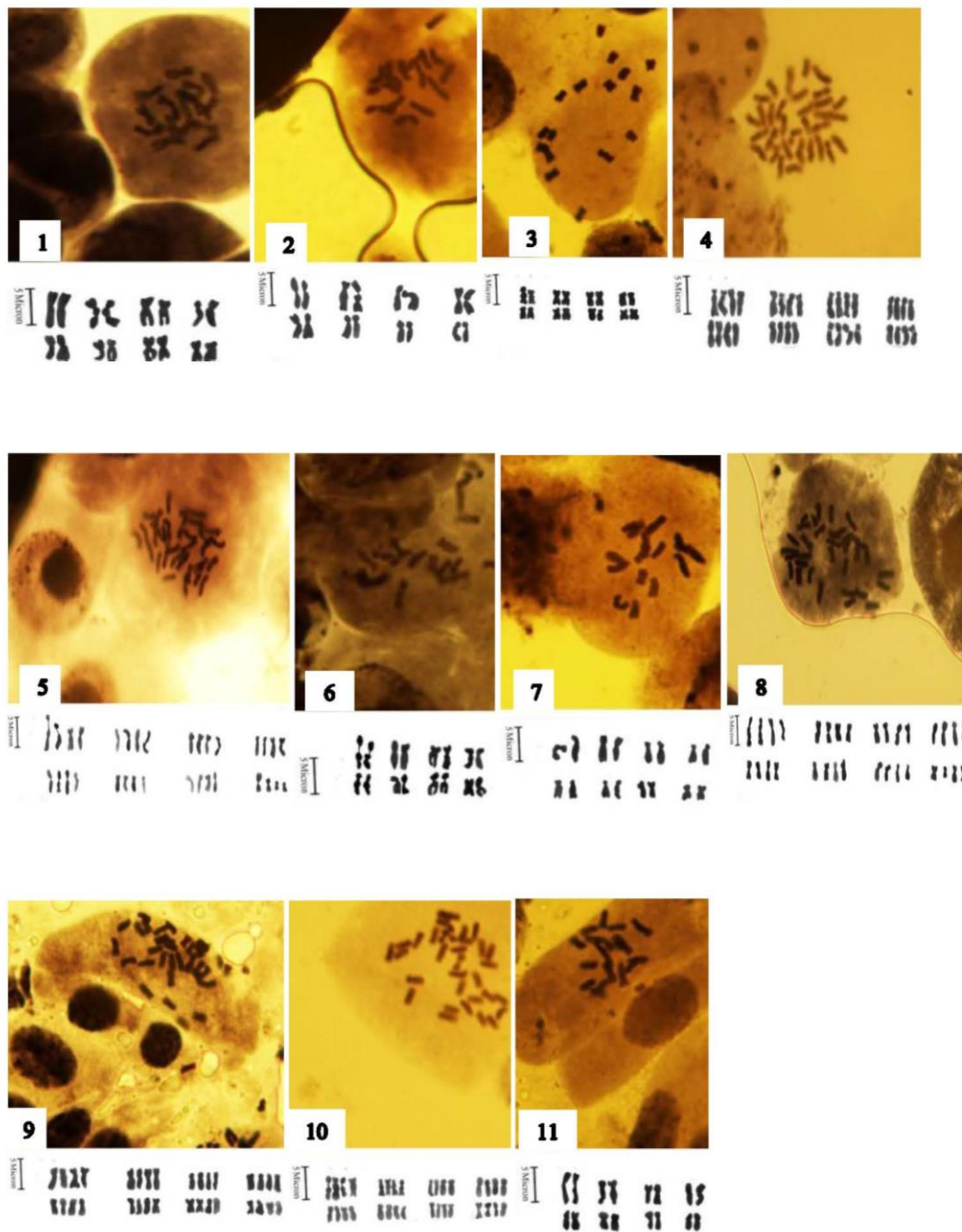


Fig. 1. Karyotypes and karyograms of 11 accessions of *Astragalus* L. 1, *Astragalus cephalanthus* DC. (105647); 2, *A. curvirostris* Boiss. (105636); 3, *A. cyclophyllus* G. Beck (107732); 4, *A. cyclophyllus* G. Beck (107734); 5, *A. cyclophyllus* G. Beck (105645); 6, *A. cyclophyllus* G. Beck (107735); 7, *A. dactylocarpus* Boiss. (105648); 8, *A. fasciculifolius* Boiss. (107733); 9, *A. fasciculifolius* Boiss. (105659); 10, *A. jacobsii* Podlech (105639); 11, *A. megalotropis* Bunge (105654).

In *A. cyclophyllus* (105645 TARI and 107735 TARI) a single pair of satellite on short arm were observed. Previous studies reported that *A. stella* Gouan, *A. victoriae* Podlech & Kirchoff, *A. densifolius* Lam and *A. uraniolimneus* Boiss. had single satellite chromosome pair (Badr & Sharawy 2007; Bozkurt & al. 2019; Gedik & al. 2019; Martin & al. 2019), and *A. effusus* Bunge had two satellite chromosome pairs (Yousefzadeh & al. 2010). On the other hand, Konichenko & al. (2014) did not observe satellites in the karyotype of *A. sericeocanus* Gontsch. The satellite represents 50-80% on the long arm, depending on the species (Dopchiz & al. 1995). Dopchiz & al. (1995) and Bozkurt & al. (2019) reported species of *Astragalus* possess satellites on the long arm of chromosome, while Badr & Sharawy (2007) and Yousefzadeh & al. (2010) found satellites were always located on the short arms. In *A. cyclophyllus* (105645 TARI), only one of the two pairs of homologous chromosomes has satellites that appear to be allotetraploid. However, more study such as Gimmsa C-bandind or genomic *in situ* hybridization (GISH) methods is required to determine their origin information, that is, whether these tetraploids are allo- or autotetraploids.

Karyotype asymmetry is one of the most important parameters in chromosomal data (Martin & al. 2019). All studied accessions show relatively symmetrical karyotypes, consisting of metacentric and submetacentric chromosomes. Recent studies also confirm similar karyotypes for *Astragalus* (Bozkurt & al. 2019; Gedik & al. 2019; Sheidai & Ghahremaninejad 2009; Sheidai & al. 2009; Yousefzadeh & al. 2010). Some studied *Astragalus* species differed in their karyotype formulae. Even, the populations belong to the same species were divided into different karyotype groups. Similarly, Konichenko & al. (2014) observed that the karyotype formulae of *A. sericeocanus* Gontsch populations were divided into three groups. This indicates that changes in the form of their chromosomes possibly due to structural changes like inversion, translocation, etc., which is considered as a qualitative change in the genome (Sheidai & Ghahremaninejad 2009).

Astragalus megalotropis Bunge showed the most symmetrical karyotype as 41.07, 0.19, 1.50 and 0.41 for TF%, A₁, AR and CI, respectively. Of all the investigated accessions, *A. dactylocarpus* Boiss. presented the most asymmetrical karyotype with a TF% of 32.37%, A₁ index of 0.49, AR of 2.16 and CI of 0.33. The A₂ index ranged from 0.12 (*A. jacobsii* Podlech) to 0.23 (*A. fasciculifolius* Boiss. (107733TARI)). Based on the results of interchromosomal asymmetry with the value of lowest A₂ (0.12) and DRL (2.33), *A. jacobsii*

Podlech possessed the least asymmetrical karyotype.

Four types of Stebbins symmetry class were found: 1A, 2A, 3A and 2B. The 2B type included *A. fasciculifolius* Boiss. (107733 TARI); 3A type included two accessions (*A. cyclophyllus* (107735 TARI) and *A. dactylocarpus* Boiss. and the remaining 8 accessions belonged to the 1A (2 accessions) and 2A (6 accessions) types. However, Stebbins classification of *Astragalus* species is generally reported in 1A or 2A classes (Sheidai & Ghahremaninejad, 2009; Sheidai & al. 2009), which indicates more asymmetry karyotypes in some accessions studied.

REFERENCES

- Badr, A. & Sharawy, Sh. M. 2007: Karyotype analysis and systematic relationships in the Egyptian *Astragalus* L. (Fabaceae). -Int. J. Bot. 3: 147-159.
- Bagheri, A., Maassoumi, A. A. & Ghahremaninejad, F. 2011: A taxonomic revision of the genus *Astragalus* L. (Fabaceae) in Zanjan province, Iran and describing a new species. -Taxon. Biosystem. 8: 7-16.
- Bozkurt, M., Ertuğrul, K. & Uysal, T. 2019: Chromosome number and karyomorphologic studies of endemic *Astragalus victoriae* Podlech & Kirchoff. -Biol. Divers. Conser. 12 (2): 169-172.
- Dopchiz, L. P., Gomez-Sosa, E. & Poggio, L. 1995: Karyotype and nuclear DNA content of six species of *Astragalus* (Leguminosae). Cytologia. 60: 329-335.
- Gedik, O., Kurşat, M. & Kiran, Y. 2019: Karyological studies on nine *Astragalus* L. taxa in Turkey. -KSU J. Agri. Nat. 22 (1): 35-44.
- Ghahremaninejad, F. 2015: Notes about *Astragalus* (Leguminosae) in Iran. -Ann. Naturhist. Mus. Wien. B. 117: 279-281.
- Goldblatt, P. & Johnson, D. E. 2010: Index to plant chromosome numbers 2004-2006. -Regnum Veg. 106: 1-242.
- ILDIS [International Legume Database & Information Service]. 2002: World Database of Legumes. ILDIS, UK.
- Konichenko, E. S., Selyutina, I. Y., Dorogina, O. V. & Sandanov, D. V. 2014: Karyotype studies endemic plant species *Astragalus sericeocanus* Gontsch. (Fabaceae) around Lake Baikal, Siberia. Caryologia. 67 (2): 172-177.
- Levan, A., Fredga, K. & Sandberg, A. A. 1964: Nomenclature for centromeric position on chromosomes. Hereditas. 52: 201-220.
- Lock, J. M. & Simpson, K. 1991: Legumes of West Asia, A Check List. Royal Botanical Gardens, UK.

- Mabberley, D. J. 1997: The plant book. Cambridge University Press, UK.
- Maassoumi, A. A. 1987: Notes on the genus *Astragalus* in Iran I, cytotaxonomic studies on some species. - Iran. J. Bot. 3: 117-128.
- Massoumi, A. A. 1998: Old World Check-List of *Astragalus*. Research Institute of Forests and Rangelands Press, Iran.
- Marhold, K. & Breitwieser, I. (eds.) 2009: IAPT/IOPB chromosome data 8. Taxon. 58: 1281-1289.
- Martin, E., İcyer Doğan, G., Karaman Erkul, S. & Eroğlu, H. E. 2019: Karyotype analyses of 25 Turkish taxa of *Astragalus* from the sections *Macrophyllum*, *Hymenostegis*, *ymenocoleus*, and *Anthylloidei* (Fabaceae). -Turk. J. Bot. 43: 232-242.
- Miri, S. M. & Shamsolshoara Y. 2019: Phytochemical and pharmacological properties of *Astragalus* species. -Proc. 2nd Nat. Conf. Med. Plant. Species Affect. Fatty Liver, 13 Jun. 2019, Tehran, Iran.
- Ranjbar, M., Hadidchi, A. & Riahi, H. 2014: Chromosome number reports in *Astragalus* sect. *Onobrychoidei* (Fabaceae) from Iran. -Taxon. Biosystem. 21: 71-82.
- Ranjbar, M., Karamian, R. & Nouri, S. 2011: Impact of cytotoxicity on meiosis in *Astragalus cyclophyllos* Beck (Fabaceae) from Iran. Caryologia. 64 (3): 256-264.
- Ranjbar, M. & Mahmoudian, B. 2015: An overview on cytogenetics of the genus *Astragalus* subgenus *Hypoglottis* (Fabaceae). Caryologia. 68 (2): 109-124.
- Sheidai, M. & Ghahremaninejad, F. 2009: New chromosome number and karyotype analysis in four *Astragalus* L. (Fabaceae) species. -Iran. J. Bot. 15 (1): 21-26.
- Sheidai, M., Zarre, Sh. & Ismeilzadeh, J. 2009: New chromosome number reports in tragacanthic *Astragalus* species. Caryologia. 62 (1): 30-36.
- Stebbins, G. L. 1971: Chromosomal Evolution in Higher Plants. Edward Arnold Press, UK.
- Yousefzadeh, K., Houshmand, S. & Zamani Dadane, G. 2010: Karyotype analysis of *Astragalus effusus* Bunge (Fabaceae). Caryologia. 63 (3): 257-261.
- Zarco, C. R. 1986: A new method for estimating karyotype asymmetry. Taxon. 35: 526-530.